SEQUENCE LISTING

```
<110> BRIGGS, Kristen
     GLANCY, Todd
     HEIN, Mitch B.
     HIATT, Andrew C.
     KARNOUP, Anton S.
     ANDERSON, W.H. Kerr
     PAREDDY, Dayakar
     PETOLINO, Joseph
     RUBIN-WILSON, Elizabeth
     TAYLOR, Doug
     Roberts, Jean L.
     The Dow Chemical Company
     Dow Agrosciences, LLC
     Epicyte Pharmaceutical, Inc.
<120> Plant production of immunoglobulins with reduced fucosylation
<130> 038136-5001-US
<150> US 60/429,385
<151>
      2002-11-27
<150> PCT/US03/037905
<151> 2003-11-28
<160> 85
<170> PatentIn version 3.1
<210> 1
<211> 1494
<212> DNA
<213> Herpes simplex virus
<220>
<221> CDS
<222>
      (1)..(1494)
<220>
<221> misc feature
<223> HSV Heavy Chain sequence
<400> 1
atg gga tgg agc tgg atc ttt ctc ttc ctc ctg tca gga gct gca ggt
                                                                       48
Met Gly Trp Ser Trp Ile Phe Leu Phe Leu Leu Ser Gly Ala Ala Gly
                                                                       96
gtc cat tgc cag gtt cag ctc gtg cag tca ggt gct gag gtg aag aag
Val His Cys Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
            20
                                                                      144
cct ggc tcc tcg gtg aag gtc tcc tgc aag gct tct gga ggt tcc ttc
Pro Gly Ser Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Ser Phe
       35
                            40
                                                45
```

								caa Gln		192
								aac Asn		240
								tcc Ser		288
								acg Thr 110		336
								gtc Val		384
								gtc Val		432
								tgt Cys		480
								ggc Gly		528
								cag Gln 190		576
								gac Asp		624
								cta Leu		672
_		_		_		_		agc Ser		720
								cca Pro		768
								agg Arg 270		816

Á

_			-		 _	_				_	-	gcg Ala	864
												acc Thr	912
		_		_	 _	_	_	-				cct Pro	 960
												ggc Gly 335	1008
												gcc Ala	1056
												ggc Gly	1104
				_	-	-	_	_	-			gag Glu	1152
												ttc Phe	1200
												ctg Leu 415	1248
												caa Gln	1296
												gag Glu	1344
	_	_	 -			-	-					gcc Ala	1392
												aaa Lys	1440
												acc Thr 495	1488
tac	tga												1494

•

Tyr

<210> 2

<211> 497

<212> PRT

<213> Herpes simplex virus

<400> 2

Met Gly Trp Ser Trp Ile Phe Leu Phe Leu Leu Ser Gly Ala Ala Gly
1 10 15

Val His Cys Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys 20 25 30

Pro Gly Ser Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Ser Phe 35 40 45

Ser Ser Tyr Ala Ile Asn Trp Val Arg Gln Ala Pro Gly Gln Gly Leu 50 55 60

Glu Trp Met Gly Gly Leu Met Pro Ile Phe Gly Thr Thr Asn Tyr Ala
65 70 75 80

Gln Lys Phe Gln Asp Arg Leu Thr Ile Thr Ala Asp Val Ser Thr Ser 85 90 95

Thr Ala Tyr Met Gln Leu Ser Gly Leu Thr Tyr Glu Asp Thr Ala Met 100 105 110

Tyr Tyr Cys Ala Arg Val Ala Tyr Met Leu Glu Pro Thr Val Thr Ala 115 120 125

Gly Gly Leu Asp Val Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser 130 140

Ala Ser Pro Thr Ser Pro Lys Val Phe Pro Leu Ser Leu Cys Ser Thr 145 150 155 160

Gln Pro Asp Gly Asn Val Val Ile Ala Cys Leu Val Gln Gly Phe Phe 165 170 175

Pro Gln Glu Pro Leu Ser Val Thr Trp Ser Glu Ser Gly Gln Gly Val 180 185 190

Thr Ala Arg Asn Phe Pro Pro Ser Gln Asp Ala Ser Gly Asp Leu Tyr Thr Thr Ser Ser Gln Leu Thr Leu Pro Ala Thr Gln Cys Leu Ala Gly Lys Ser Val Thr Cys His Val Lys His Tyr Thr Asn Pro Ser Gln Asp Val Thr Val Pro Cys Pro Val Pro Ser Thr Pro Pro Thr Pro Ser Pro Ser Thr Pro Pro Thr Pro Ser Pro Ser Cys Cys His Pro Arg Leu Ser Leu His Arg Pro Ala Leu Glu Asp Leu Leu Gly Ser Glu Ala Asn Leu Thr Cys Thr Leu Thr Gly Leu Arg Asp Ala Ser Gly Val Thr Phe Thr Trp Thr Pro Ser Ser Gly Lys Ser Ala Val Gln Gly Pro Pro Glu Arg Asp Leu Cys Gly Cys Tyr Ser Val Ser Ser Val Leu Pro Gly Cys Ala Glu Pro Trp Asn His Gly Lys Thr Phe Thr Cys Thr Ala Ala Tyr Pro Glu Ser Lys Thr Pro Leu Thr Ala Thr Leu Ser Lys Ser Gly Asn Thr Phe Arg Pro Glu Val His Leu Leu Pro Pro Pro Ser Glu Glu Leu Ala Leu Asn Glu Leu Val Thr Leu Thr Cys Leu Ala Arg Gly Phe Ser Pro Lys Asp Val Leu Val Arg Trp Leu Gln Gly Ser Gln Glu Leu Pro

Arg Glu Lys Tyr Leu Thr Trp Ala Ser Arg Gln Glu Pro Ser Gln Gly 420 425 Thr Thr Thr Phe Ala Val Thr Ser Ile Leu Arg Val Ala Ala Glu Asp 440 Trp Lys Lys Gly Asp Thr Phe Ser Cys Met Val Gly His Glu Ala Leu 450 455 Pro Leu Ala Phe Thr Gln Lys Thr Ile Asp Arg Leu Ala Gly Lys Pro 465 470 475 Thr His Val Asn Val Ser Val Val Met Ala Glu Val Asp Gly Thr Cys 485 Tyr <210> 3 <211> 57 <212> DNA <213> Artificial sequence <220> <223> Heavy chain signal peptide <220> <221> CDS <222> (1)..(57) <400> 3 atg gga tgg agc tgg atc ttt ctc ttc ctc ctg tca gga gct gca ggt 48 Met Gly Trp Ser Trp Ile Phe Leu Phe Leu Leu Ser Gly Ala Ala Gly 1 5 10 15 57 gtc cat tgc Val His Cys <210> 4 <211> 19 <212> PRT <213> Artificial sequence <223> Heavy chain signal peptide <400> 4

Met Gly Trp Ser Trp Ile Phe Leu Phe Leu Leu Ser Gly Ala Ala Gly 1 5 10 15

Val His Cys

<210> 5 <211> 1368 <212> DNA <213> Artificial sequence														
Mature heavy chain sequence														
<220> <221> CDS <222> (1)(1368)														
<pre><400> 5 cag gtt cag ctc gtg cag tca ggt gct gag gtg aag aag cct ggc tcc Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser 1</pre>	8													
tcg gtg aag gtc tcc tgc aag gct tct gga ggt tcc ttc agc tcc tat Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Ser Phe Ser Ser Tyr 20 25 30	6													
gct atc aac tgg gtg agg caa gct cct gga caa ggg ctt gag tgg atg Ala Ile Asn Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met 35 40 45	4													
gga ggg ctc atg cct atc ttt ggg aca aca aac tac gcg cag aag ttc Gly Gly Leu Met Pro Ile Phe Gly Thr Thr Asn Tyr Ala Gln Lys Phe 50 55 60	2													
cag gac agg ctc acg att acc gcg gac gta tcc acg agt aca gcc tac Gln Asp Arg Leu Thr Ile Thr Ala Asp Val Ser Thr Ser Thr Ala Tyr 65 70 75 80	0													
atg caa ctg agc ggc ctg aca tat gaa gac acg gcc atg tat tac tgt Met Gln Leu Ser Gly Leu Thr Tyr Glu Asp Thr Ala Met Tyr Tyr Cys 85 90 95	8													
gcg aga gtt gcc tac atg ctt gaa cct acc gtc act gca ggt ggt ttg Ala Arg Val Ala Tyr Met Leu Glu Pro Thr Val Thr Ala Gly Gly Leu 100 105 110	6													
gac gtc tgg ggc caa ggg acc ttg gtc acc gtc tcc tcc gca tcc ccg Asp Val Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Pro 115 120 125	4													
acc agc ccg aag.gtc ttc ccg ctg agc ctc tgt agc acc cag cca gat Thr Ser Pro Lys Val Phe Pro Leu Ser Leu Cys Ser Thr Gln Pro Asp 130 135 140	2													

.

					gcc Ala 150											480
					tgg Trp											528
					cag Gln											576
_	_	_			ccg Pro	-		_	-				_			624
					cac His											672
	_		_		tca Ser 230								-			720
					tca Ser	_	_				_		_			768
					ctg Leu											816
					aga Arg											864
					agc Ser											912
					gtg Val 310											960
				_	acc Thr			-		-	-			-	_	1008
					gcc Ala											1056
		-		_	ctg Leu	_	_	_	_							1104
gag	ctg	gtg	acg	ctg	acg	tgc	ctg	gcg	cgc	ggc	ttc	agc	ccc	aag	gac	1152

.

Glu	Leu 370	Val	Thr	Leu	Thr	Cys 375	Leu	Ala	Arg	Gly	Phe 380	Ser	Pro	Lys	Asp	
				tgg Trp												1200
				gca Ala 405												1248
				tcg Ser												1296
				tcc Ser												1344
				acc Thr												1368
<213 <220 <223	2> I 3> <i>I</i> 0>	Artif Matur		al se eavy	_		equer	nce								
			Leu	Val 5	Gln	Ser	Gly	Ala	Glu 10	Val	Lys	Lys	Pro	Gly 15	Ser	
Ser	Val	Lys	Val 20	Ser	Cys	Lys	Ala	Ser 25	Gly	Gly	Ser	Phe	Ser 30	Ser	Tyr	
Ala	Ile	Asn 35	Trp	Val	Arg	Gln	Ala 40	Pro	Gly	Gln	Gly	Leu 45	Glu	Trp	Met	
Gly	Gly 50	Leu	Met	Pro	Ile	Phe 55	Gly	Thr	Thr	Asn	Tyr 60	Ala	Gln	Lys	Phe	
Gln 65	Asp	Arg	Leu	Thr	Ile 70	Thr	Ala	Asp	Val	Ser 75	Thr	Ser	Thr	Ala	Tyr 80	

Ala Arq Val Ala Tyr Met Leu Glu Pro Thr Val Thr Ala Gly Gly Leu 105 100 Asp Val Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Pro 120 Thr Ser Pro Lys Val Phe Pro Leu Ser Leu Cys Ser Thr Gln Pro Asp 130 135 Gly Asn Val Val Ile Ala Cys Leu Val Gln Gly Phe Phe Pro Gln Glu 155 150 145 Pro Leu Ser Val Thr Trp Ser Glu Ser Gly Gln Gly Val Thr Ala Arg 170 Asn Phe Pro Pro Ser Gln Asp Ala Ser Gly Asp Leu Tyr Thr Thr Ser 185 Ser Gln Leu Thr Leu Pro Ala Thr Gln Cys Leu Ala Gly Lys Ser Val 195 200 Thr Cys His Val Lys His Tyr Thr Asn Pro Ser Gln Asp Val Thr Val 210 215 Pro Cys Pro Val Pro Ser Thr Pro Pro Thr Pro Ser Pro Ser Thr Pro 225 230 235 Pro Thr Pro Ser Pro Ser Cys Cys His Pro Arg Leu Ser Leu His Arg 245 250 Pro Ala Leu Glu Asp Leu Leu Gly Ser Glu Ala Asn Leu Thr Cys 260 265 Thr Leu Thr Gly Leu Arg Asp Ala Ser Gly Val Thr Phe Thr Trp Thr Pro Ser Ser Gly Lys Ser Ala Val Gln Gly Pro Pro Glu Arg Asp Leu 290 295 Cys Gly Cys Tyr Ser Val Ser Ser Val Leu Pro Gly Cys Ala Glu Pro 310 305 315

Trp Asn His Gly Lys Thr Phe Thr Cys Thr Ala Ala Tyr Pro Glu Ser 325 330 Lys Thr Pro Leu Thr Ala Thr Leu Ser Lys Ser Gly Asn Thr Phe Arg Pro Glu Val His Leu Leu Pro Pro Pro Ser Glu Glu Leu Ala Leu Asn Glu Leu Val Thr Leu Thr Cys Leu Ala Arg Gly Phe Ser Pro Lys Asp 370 375 Val Leu Val Arg Trp Leu Gln Gly Ser Gln Glu Leu Pro Arg Glu Lys 390 395 Tyr Leu Thr Trp Ala Ser Arg Gln Glu Pro Ser Gln Gly Thr Thr 405 410 Phe Ala Val Thr Ser Ile Leu Arg Val Ala Ala Glu Asp Trp Lys Lys 420 425 430 Gly Asp Thr Phe Ser Cys Met Val Gly His Glu Ala Leu Pro Leu Ala 435 Phe Thr Gln Lys Thr Ile Asp Arg 450 455 <210> 7 <211> 69 <212> DNA <213> Artificial sequence <220> <223> heavy chain tailpiece <220> <221> CDS <222> (1)..(69) <400> 7 ttg gcg ggt aaa ccc acc cat gtc aat gtg tct gtc atg gcg gag Leu Ala Gly Lys Pro Thr His Val Asn Val Ser Val Val Met Ala Glu 15 5 10 gtg gac ggc acc tgc tac tga 69 Val Asp Gly Thr Cys Tyr

20

```
<210> 8
<211> 22
<212> PRT
<213> Artificial sequence
<220>
<223>
      heavy chain tailpiece
<400>
Leu Ala Gly Lys Pro Thr His Val Asn Val Ser Val Val Met Ala Glu
Val Asp Gly Thr Cys Tyr
<210> 9
<211> 702
<212> DNA
<213> Herpes simplex virus
<220>
<221> CDS
<222> (1)..(702)
<220>
<221> misc feature
<223> HSV light chain sequence
<400> 9
atg gga tgg tcc tgg atc ttt ctc ttc ctt ctg tca gga gct gca ggt
                                                                       48
Met Gly Trp Ser Trp Ile Phe Leu Phe Leu Leu Ser Gly Ala Ala Gly
gtc cac tgc gag atc gtg ctc acg cag tct cca ggc acc ctg tct ttg
                                                                       96
Val His Cys Glu Ile Val Leu Thr Gln Ser Pro Gly Thr Leu Ser Leu
            20
                                25
                                                    30
tcg cca ggg gaa cgt gcc acc ctc tcc tgc cgg gcc agt cag tcc gtt
                                                                      144
Ser Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val
                            40
                                                                      192
tee age geg tae ett gee tgg tae eag eag aag eet gge eaa get eee
Ser Ser Ala Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro
                                                                      240
agg etc etc atc tat ggt geg tec age agg get act gge att eca gae
Arg Leu Leu Ile Tyr Gly Ala Ser Ser Arg Ala Thr Gly Ile Pro Asp
                                                                      288
cgc ttc tca ggc agt ggg tct ggg aca gac ttc acg ctc acc att agc
Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser
                85
                                    90
```

agg ctg gaa cct gag Arg Leu Glu Pro Glu 100				336
cgc tca ccc acg ttc Arg Ser Pro Thr Phe 115				384
gtg gct gca ccg tcg Val Ala Ala Pro Ser 130			Asp Glu Gln Leu	432
aag tot ggc act gcc Lys Ser Gly Thr Ala 145			*	480
aga gag gcg aag gta Arg Glu Ala Lys Val 165	Gln Trp Lys			528
aac tcc caa gag tcc Asn Ser Gln Glu Ser 180			9 9	576
agc ctc agc aac acc Ser Leu Ser Asn Thr 195			• • • • • • • • • • • • • • • • • • • •	624
aag gtc tac gcc tgc Lys Val Tyr Ala Cys 210			Arg Ser Pro Val	672
aca aag agc ttc aac Thr Lys Ser Phe Asn 225		_	•	702
<210> 10 <211> 233 <212> PRT <213> Herpes simpl	ex virus		•	
<400> 10				
Met Gly Trp Ser Trp 1 5	Ile Phe Leu	Phe Leu Leu Ser 10	Gly Ala Ala Gly 15	
Val His Cys Glu Ile 20	Val Leu Thr	Gln Ser Pro Gly 25	Thr Leu Ser Leu 30	
Ser Pro Gly Glu Arg 35	Ala Thr Leu 40	Ser Cys Arg Ala	Ser Gln Ser Val 45	

Ser Ser Ala Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro 50 55 60

Arg Leu Leu Ile Tyr Gly Ala Ser Ser Arg Ala Thr Gly Ile Pro Asp 65 70 75 80

Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser 85 90 95

Arg Leu Glu Pro Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Gly
100 105 110

Arg Ser Pro Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr 115 120 125

Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu 130 135 140

Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro 145 $$\sim150 155 160

Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly
165 170 175

Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr 180 185 190

Ser Leu Ser Asn Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His 195 200 205

Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Arg Ser Pro Val 210 215 220

Thr Lys Ser Phe Asn Arg Gly Glu Cys 225 230

<210> 11

<211> 57

<212> DNA

<213> Artificial sequence

<220>

<223> Light chain signal peptide

<220>

```
<221> CDS
<222>
      (1)..(57)
<400> 11
                                                                       48
atg gga tgg tcc tgg atc ttt ctc ttc ctt ctg tca gga gct gca ggt
Met Gly Trp Ser Trp Ile Phe Leu Phe Leu Leu Ser Gly Ala Ala Gly
                                    10
                                                                       57
gtc cac tgc
Val His Cys
<210> 12
<211> 19
<212> PRT
<213> Artificial sequence
<220>
<223> Light chain signal peptide
<400> 12
Met Gly Trp Ser Trp Ile Phe Leu Phe Leu Leu Ser Gly Ala Ala Gly
Val His Cys
<210> 13
<211> 642
<212> DNA
<213> Artificial sequence
<220>
<223> Mature light chain sequence
<220>
<221> CDS
<222> (1)..(642)
<400> 13
                                                                       48
gag atc gtg ctc acg cag tct cca ggc acc ctg tct ttg tcg cca ggg
Glu Ile Val Leu Thr Gln Ser Pro Gly Thr Leu Ser Leu Ser Pro Gly
                5
                                                                       96
gaa cgt gcc acc ctc tcc tgc cgg gcc agt cag tcc gtt tcc agc gcg
Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Ser Ala
                                                                      144
tac ctt gcc tgg tac cag cag aag cct ggc caa gct ccc agg ctc ctc
Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu
                                                                      192
atc tat ggt gcg tcc agc agg gct act ggc att cca gac cgc ttc tca
```

Ile	Tyr 50	Gly	Ala	Ser	Ser	Arg 55	Ala	Thr	Gly	Ile	Pro 60	Asp	Arg	Phe	Ser	
		ggg Gly														240
		gat Asp														288
		ggc Gly														336
_	_	gtc Val 115				_			-		_	_				384
	-	tct Ser	-		_	_	_					_				432
		cag Gln														480
		gtt Val			_	_	-	-	_	_						528
		ttg Leu														576
		gaa Glu 195														624
		cgg Arg														642
<21 <21 <21 <21	1> 2 2> 1	14 214 PRT Arti	ficia	al se	equer	nce										
<22 <22		Matui	re l	ight	cha	in se	equei	nce								
<40	0> :	14														
Glu 1	Ile	Val	Leu	Thr 5	Gln	Ser	Pro	Gly	Thr 10	Leu	Ser	Leu	Ser	Pro 15	Gly	

Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Ser Ala 20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu 35 40 45

Ile Tyr Gly Ala Ser Ser Arg Ala Thr Gly Ile Pro Asp Arg Phe Ser 50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Arg Leu Glu 65 70 75 80

Pro Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Gly Arg Ser Pro 85 90 95

Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala Ala 100 105 110

Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly 115 120 125

Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala 130 135 140

Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser Gln 145 150 155 160

Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu Ser 165 170 175

Asn Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Val Tyr 180 185 190

Ala Cys Glu Val Thr His Gln Gly Leu Arg Ser Pro Val Thr Lys Ser 195 200 205

Phe Asn Arg Gly Glu Cys 210

<210> 15

<211> 9144

<212> DNA

<213> Artificial sequence

<220> <223> pDAB635 (ubiH) sequence

<400> 15 60 tcgcgcgttt cggtgatgac ggtgaaaacc tctgacacat gcagctcccg gagacggtca cagettgtct gtaageggat geegggagea gacaageeeg teagegggtg 120 ttggcgggtg tcggggctgg cttaactatg cggcatcaga gcagattgta ctgagagtgc 180 accatatgcg gtgtgaaata ccgcacagat gcgtaaggag aaaataccgc atcaggcgcc 240 attcgccatt caggctgcgc aactgttggg aagggcgatc ggtgcgggcc tcttcgctat 300 tacgccagct ggcgaaaggg ggatgtgctg caaggcgatt aagttgggta acgccagggt 360 420 tttcccagtc acgacgttgt aaaacgacgg ccagtgaatt acaccggtgt gatcatgggc 480 cgcgattaaa aatcccaatt atatttggtc taatttagtt tggtattgag taaaacaaat 540 tcgaaccaaa ccaaaatata aatatatagt ttttatatat atgcctttaa gactttttat 600 agaattttct ttaaaaaata tctagaaata tttgcgactc ttctggcatg taatatttcg 660 ttaaatatga agtgctccat ttttattaac tttaaataat tggttgtacg atcactttct 720 tatcaagtgt tactaaaatg cgtcaatctc tttgttcttc catattcata tgtcaaaatc 780 tatcaaaatt cttatatatc tttttcgaat ttgaagtgaa atttcgataa tttaaaaatta 840 aatagaacat atcattattt aggtatcata ttgattttta tacttaatta ctaaatttgg 900 ttaactttga aagtgtacat caacgaaaaa ttagtcaaac gactaaaata aataaatatc 960 atgtgttatt aagaaaattc tcctataaga atattttaat agatcatatg tttgtaaaaa 1020 aaattaattt ttactaacac atatatttac ttatcaaaaa tttgacaaag taagattaaa 1080 ataatattca tctaacaaaa aaaaaaccag aaaatgctga aaacccggca aaaccgaacc 1140 aatccaaacc gatatagttg gtttggtttg attttgatat aaaccgaacc aactcggtcc atttgcaccc ctaatcataa tagctttaat atttcaagat attattaagt taacgttgtc 1200 1260 aatatcctgg aaattttgca aaatgaatca agcctatatg gctgtaatat gaatttaaaa 1320 gcagctcgat gtggtggtaa tatgtaattt acttgattct aaaaaaatat cccaagtatt 1380 aataatttct gctaggaaga aggttagcta cgatttacag caaagccaga atacaaagaa 1440 ccataaagtg attgaagctc gaaatatacg aaggaacaaa tatttttaaa aaaatacgca atgacttgga acaaaagaaa gtgatatatt ttttgttctt aaacaagcat cccctctaaa 1500 1560 gaatggcagt tttcctttgc atgtaactat tatgctccct tcgttacaaa aattttggac

1620 tactattggg aacttettet gaaaatagtg gecacegett aattaacace ggtggeeegg 1680 gcaagcggcc gcattcccgg gaagctaggc caccgtggcc cgcctgcagg ggaagcttgc 1740 atgcctgcag atccccgggg atcctctaga gtcgacctgc agtgcagcgt gacccggtcg tgcccctctc tagagataat gagcattgca tgtctaagtt ataaaaaatt accacatatt 1800 1860 ttttttgtca cacttgtttg aagtgcagtt tatctatctt tatacatata tttaaacttt aatctacgaa taatataatc tatagtacta caataatatc agtgttttag agaatcatat 1920 aaatgaacag ttagacatgg tctaaaggac aattgagtat tttgacaaca ggactctaca 1980 2040 gttttatctt tttagtgtgc atgtgttctc cttttttttt gcaaatagct tcacctatat 2100 aatacttcat ccattttatt agtacatcca tttagggttt agggttaatg gtttttatag 2160 actaattttt ttagtacatc tattttattc tattttagcc tctaaattaa gaaaactaaa 2220 actctatttt agttttttta tttaataatt tagatataaa atagaataaa ataaagtgac 2280 taaaaaattaa acaaataccc tttaagaaat taaaaaaact aaggaaacat ttttcttgtt 2340 tegagtagat aatgeeagee tgttaaaege egtegaegag tetaaeggae aecaaecage gaaccagcag cgtcgcgtcg ggccaagcga agcagacggc acggcatctc tgtcgctgcc 2400 2460 tetggacece tetegagagt teegeteeae egttggaett geteegetgt eggeateeag 2520 2580 acggcacggc agctacgggg gattcctttc ccaccgctcc ttcgctttcc cttcctcgcc 2640 cgccgtaata aatagacacc ccctccacac cctctttccc caacctcgtg ttgttcggag 2700 cgcacacaca cacaaccaga tctcccccaa atccacccgt cggcacctcc gcttcaaggt acgccgctcg tectecece ecceectet etacettete tagateggeg tteeggteea 2760 tgcatggtta gggcccggta gttctacttc tgttcatgtt tgtgttagat ccgtgtttgt 2820 gttagatccg tgctgctagc gttcgtacac ggatgcgacc tgtacgtcag acacgttctg 2880 2940 attgctaact tgccagtgtt tctctttggg gaatcctggg atggctctag ccgttccgca 3000 gacgggatcg atttcatgat tttttttgtt tcgttgcata gggtttggtt tgcccttttc 3060 ctttatttca atatatgccg tgcacttgtt tgtcgggtca tcttttcatg cttttttttg 3120 tettggttgt gatgatgtgg tetggttggg eggtegttet agateggagt agaattetgt 3180 ttcaaactac ctggtggatt tattaatttt ggatctgtat gtgtgtgcca tacatattca 3240 tagttacgaa ttgaagatga tggatggaaa tatcgatcta ggataggtat acatgttgat 3300 gcgggtttta ctgatgcata tacagagatg ctttttgttc gcttggttgt gatgatgtgg

3360 tgtggttggg cggtcgttca ttcgttctag atcggagtag aatactgttt caaactacct 3420 ggtgtattta ttaattttgg aactgtatgt gtgtgtcata catcttcata gttacgagtt 3480 taagatggat ggaaatatcg atctaggata ggtatacatg ttgatgtggg ttttactgat 3540 gcatatacat gatggcatat gcagcatcta ttcatatgct ctaaccttga gtacctatct attataataa acaagtatgt tttataatta ttttgatctt gatatacttg gatgatggca 3600 tatgcagcag ctatatgtgg attitttag ccctgccttc atacgctatt tatttgcttg 3660 gtactgtttc ttttgtcgat gctcaccctg ttgtttggtg ttacttctgc agggtacccc 3720 3780 eggggtegae catggecaae aageaeetga geeteteeet etteetegtg eteeteggee 3840 totocgooto cotogocago ggocaggtto agotogtgoa gtoaggggot gaggtgaaga 3900 agcctgggtc ctcggtgaag gtctcctgca aggcttctgg aggttccttc agcagctatg 3960 ctatcaactg ggtgcgacag gcccctggac aagggcttga gtggatggga gggctcatgc 4020 ctatctttgg gacaacaaac tacgcacaga agttccagga cagactcacg attaccgcgg 4080 acgtatccac gagtacagcc tacatgcagc tgagcggcct gacatatgaa gacacggcca 4140 tgtattactg tgcgagagtt gcctatatgt tggaacctac cgtcactgca gggggtttgg 4200 acgtctgggg caaagggacc acggtcaccg tctccccagc atccccgacc agccccaagg 4260 tetteceget gageetetge ageacceage cagatgggaa egtggteate geetgeetgg 4320 tocagggott ottoccocag gagocactoa gtgtgacotg gagogaaago ggacagggog 4380 tgaccgccag aaacttccca cccagccagg atgcctccgg ggacctgtac accacgagca 4440 gecagetgae cetgeeggee acacagtgee tageeggeaa gteegtgaea tgecaegtga 4500 agcactacac gaatcccagc caggatgtga ctgtgccctg cccagttccc tcaactccac ctaccccatc teceteaact ecacetacce catetecete atgetgeeac eccegactgt 4560 cactgcaccg accggccctc gaggacctgc tcttaggttc agaagcgaac ctcacgtgca 4620 cactgaccgg cctgagagat gcctcaggtg tcaccttcac ctggacgccc tcaagtggga 4680 4740 agagegetgt teaaggaeea eetgagegtg acetetgtgg etgetaeage gtgteeagtg 4800 tectgeeggg etgtgeegag eettggaate atgggaagae etteaettge aetgetgeet 4860 accecgagte caagaceceg etaacegeea eceteteaaa ateeggaaae acatteegge 4920 ccgaggtcca cctgctgccg ccgccgtcgg aggagctggc cctgaacgag ctggtgacgc 4980 tgacgtgcct ggcacgtggc ttcagcccca aggacgtgct ggttcgctgg ctgcaggggt

5040 cacaggaget gccccgcgag aagtacctga cttgggcatc ccggcaggag cccagccagg 5100 gcaccaccac cttcgctgtg accagcatac tgcgcgtggc agccgaggac tggaagaagg 5160 gggacacctt ctcctgcatg gtgggccacg aggccctgcc gctggccttc acacagaaga 5220 ccatcgaccg cttggcgggt aaacccaccc atgtcaatgt gtctgttgtc atggcggagg 5280 tggacggcac ctgctactga gttaaactga gggcactgaa gtcgcttgat gtgctgaatt gtttgtgatg ttggtggcgt attttgttta aataagtaag catggctgtg attttatcat 5340 atgatcgatc tttggggttt tatttaacac attgtaaaat gtgtatctat taataactca 5400 atgtataaga tgtgttcatt cttcggttgc catagatctg cttatttgac ctgtgatgtt 5460 5520 ttgactccaa aaaccaaaat cacaactcaa taaactcatg gaatatgtcc acctgtttct tgaagagttc atctaccatt ccagttggca tttatcagtg ttgcagcggc gctgtgcttt 5580 5640 gtaacataac aattgttacg gcatatatcc aacggccggc ctaggccacg gtggccagat 5700 ccactagttc tagagcggcc gcttaattaa atttaaatgt ttaaactagg cctcctgcag 5760 ggtttaaact tgccgtggcc tattttcaga agaagttccc aatagtagtc caaaattttt gtaacgaagg gagcataata gttacatgca aaggaaaact gccattcttt agaggggatg 5820 5880 cttgtttaag aacaaaaat atatcacttt cttttgttcc aagtcattgc gtatttttt 5940 aaaaatattt gttccttcgt atatttcgag cttcaatcac tttatggttc tttgtattct 6000 ggctttgctg taaatcgtag ctaaccttct tcctagcaga aattattaat acttgggata 6060 tttttttaga atcaagtaaa ttacatatta ccaccacatc gagctgcttt taaattcata 6120 ttacagccat ataggcttga ttcattttgc aaaatttcca ggatattgac aacgttaact 6180 taataatatc ttgaaatatt aaagctatta tgattagggg tgcaaatgga ccgagttggt teggtttata teaaaateaa accaaaceaa etatateggt ttggattggt teggttttge 6240 cgggttttca gcattttctg gttttttttt tgttagatga atattatttt aatcttactt 6300 6360 tgtcaaattt ttgataagta aatatatgtg ttagtaaaaa ttaatttttt ttacaaacat 6420 atgatctatt aaaatattct tataggagaa ttttcttaat aacacatgat atttatttat 6480 tttagtcgtt tgactaattt ttcgttgatg tacactttca aagttaacca aatttagtaa 6540 ttaagtataa aaatcaatat gatacctaaa taatgatatg ttctatttaa ttttaaatta 6600 tcgaaatttc acttcaaatt cgaaaaagat atataagaat tttgatagat tttgacatat gaatatggaa gaacaaagag attgacgcat tttagtaaca cttgataaga aagtgatcgt 6660 6720 acaaccaatt atttaaagtt aataaaaatg gagcacttca tatttaacga aatattacat

6780 gccagaagag tcgcaaatat ttctagatat tttttaaaga aaattctata aaaagtctta 6840 aaggcatata tataaaaact atatatttat attttggttt ggttcgaatt tgttttactc 6900 aataccaaac taaattagac caaatataat tgggattttt aatcgcggcc cactagtcac 6960 cggtgtgctt ggcgtaatca tggtcatagc tgtttcctgt gtgaaattgt tatccgctca caattccaca caacatacga gccggaagca taaagtgtaa agcctggggt gcctaatgag 7020 tgagctaact cacattaatt gegttgeget cactgeeege tttecagteg ggaaacetgt 7080 cgtgccagct gcattaatga atcggccaac gcgcggggag aggcggtttg cgtattgggc 7140 7200 getetteege tteetegete actgaetege tgegeteggt egtteggetg eggegagegg 7260 tatcagetea eteaaaggeg gtaataeggt tateeaeaga ateaggggat aaegeaggaa 7320 agaacatgtg agcaaaaggc cagcaaaagg ccaggaaccg taaaaaggcc gcgttgctgg 7380 cgtttttcca taggctccgc cccctgacg agcatcacaa aaatcgacgc tcaagtcaga ggtggcgaaa cccgacagga ctataaagat accaggcgtt tccccctgga agctccctcg 7440 7500 tgcgctctcc tgttccgacc ctgccgctta ccggatacct gtccgccttt ctcccttcgg 7560 gaagcgtggc gctttctcat agctcacgct gtaggtatct cagttcggtg taggtcgttc 7620 gctccaagct gggctgtgtg cacgaacccc ccgttcagcc cgaccgctgc gccttatccg 7680 gtaactatcg tettgagtee aacceggtaa gacacgactt ategecactg geageageea 7740 ctggtaacag gattagcaga gcgaggtatg taggcggtgc tacagagttc ttgaagtggt 7800 ggcctaacta cggctacact agaaggacag tatttggtat ctgcgctctg ctgaagccag 7860 ttaccttcgg aaaaagagtt ggtagctctt gatccggcaa acaaaccacc gctggtagcg 7920 gtggtttttt tgtttgcaag cagcagatta cgcgcagaaa aaaaggatct caagaagatc 7980 ctttgatctt ttctacgggg tctgacgctc agtggaacga aaactcacgt taagggattt tggtcatgag attatcaaaa aggatcttca cctagatcct tttaaattaa aaatgaagtt 8040 8100 ttaaatcaat ctaaagtata tatgagtaaa cttggtctga cagttaccaa tgcttaatca gtgaggcacc tatctcagcg atctgtctat ttcgttcatc catagttgcc tgactccccg 8160 8220 tegtgtagat aactaegata egggaggget taccatetgg eeceagtget geaatgatae cgcgagaccc acgctcaccg gctccagatt tatcagcaat aaaccagcca gccggaaggg 8280 8340 ccgagcgcag aagtggtcct gcaactttat ccgcctccat ccagtctatt aattgttgcc 8400 gggaagctag agtaagtagt tegecagtta atagtttgeg caaegttgtt gecattgeta

caggcatcgt ggtgt	cacge tegtegttte	gtatggcttc	attcagctcc	ggttcccaac	8460
gatcaaggcg agtta	acatga tececeatgt	tgtgcaaaaa	agcggttagc	tccttcggtc	8520
ctccgatcgt tgtca	agaagt aagttggcco	cagtgttatc	actcatggtt	atggcagcac	8580
tgcataattc tctta	actgtc atgccatcco	, taagatgctt	ttctgtgact	ggtgagtact	8640
caaccaagtc attct	gagaa tagtgtatgo	ggcgaccgag	ttgctcttgc	ccggcgtcaa	8700
tacgggataa taccg	gegeca catageagaa	ctttaaaagt	gctcatcatt	ggaaaacgtt	8760
cttcggggcg aaaac	ctctca aggatcttad	: cgctgttgag	atccagttcg	atgtaaccca	8820
ctcgtgcacc caact	gatct tcagcatctt	ttactttcac	cagcgtttct	gggtgagcaa	8880
aaacaggaag gcaaa	aatgee geaaaaaag	gaataagggc	gacacggaaa	tgttgaatac	8940
tcatactctt ccttt	ttcaa tattattgaa	gcatttatca	gggttattgt	ctcatgagcg	9000
gatacatatt tgaat	gtatt tagaaaaata	aacaaatagg	ggttccgcgc	acatttcccc	9060
gaaaagtgcc acctg	gacgtc taagaaacca	ttattatcat	gacattaacc	tataaaaata	9120
ggcgtatcac gaggc	ecettt egte				9144

<210> 16 <211> 8352

<212> DNA <213> Artificial sequence

<220>

<223> pDAB636 (ubiL) sequence

<400> 16

60 tcgcgcgttt cggtgatgac ggtgaaaacc tctgacacat gcagctcccg gagacggtca cagettgtet gtaageggat geegggagea gacaageeeg teagggegeg teagegggtg 120 ttggcgggtg tcggggctgg cttaactatg cggcatcaga gcagattgta ctgagagtgc 180 accatatgcg gtgtgaaata ccgcacagat gcgtaaggag aaaataccgc atcaggcgcc 240 300 attegecatt caggetgege aactgttggg aagggegate ggtgegggee tettegetat 360 tacgccaget ggcgaaaggg ggatgtgctg caaggcgatt aagttgggta acgccagggt 420 tttcccagtc acgacgttgt aaaacgacgg ccagtgaatt acaccggtgt gatcatgggc 480 cgcgattaaa aatcccaatt atatttggtc taatttagtt tggtattgag taaaacaaat tcgaaccaaa ccaaaatata aatatatagt ttttatatat atgcctttaa gactttttat 540 600 agaattttct ttaaaaaata tctagaaata tttgcgactc ttctggcatg taatatttcg 660 ttaaatatga agtgctccat ttttattaac tttaaataat tggttgtacg atcactttct

tatcaagtgt	tactaaaatg	cgtcaatctc	tttgttcttc	catattcata	tgtcaaaatc	720
tatcaaaatt	cttatatatc	tttttcgaat	ttgaagtgaa	atttcgataa	tttaaaatta	780
aatagaacat	atcattattt	aggtatcata	ttgattttta	tacttaatta	ctaaatttgg	840
ttaactttga	aagtgtacat	caacgaaaaa	ttagtcaaac	gactaaaata	aataaatatc	900
atgtgttatt	aagaaaattc	tcctataaga	atattttaat	agatcatatg	tttgtaaaaa	960
aaattaattt	ttactaacac	atatatttac	ttatcaaaaa	tttgacaaag	taagattaaa	1020
ataatattca	tctaacaaaa	aaaaaaccag	aaaatgctga	aaacccggca	aaaccgaacc	1080
aatccaaacc	gatatagttg	gtttggtttg	attttgatat	aaaccgaacc	aactcggtcc	1140
atttgcaccc	ctaatcataa	tagctttaat	atttcaagat	attattaagt	taacgttgtc	1200
aatatcctgg	aaattttgca	aaatgaatca	agcctatatg	gctgtaatat	gaatttaaaa	1260
gcagctcgat	gtggtggtaa	tatgtaattt	acttgattct	aaaaaaatat	cccaagtatt	1320
aataatttct	gctaggaaga	aggttagcta	cgatttacag	caaagccaga	atacaaagaa	1380
ccataaagtg	attgaagctc	gaaatatacg	aaggaacaaa	tatttttaaa	aaaatacgca	1440
atgacttgga	acaaaagaaa	gtgatatatt	ttttgttctt	aaacaagcat	cccctctaaa	1500
gaatggcagt	tttcctttgc	atgtaactat	tatgctccct	tcgttacaaa	aattttggac	1560
tactattggg	aacttcttct	gaaaatagtg	gccaccgctt	aattaacacc	ggtggcccgg	1620
gcaagcggcc	gcattcccgg	gaagctaggc	caccgtggcc	cgcctgcagg	ggaagcttgc	1680
atgcctgcag	atccccgggg	atcctctaga	gtcgacctgc	agtgcagcgt	gacccggtcg	1740
tgcccctctc	tagagataat	gagcattgca	tgtctaagtt	ataaaaaatt	accacatatt	1800
ttttttgtca	cacttgtttg	aagtgcagtt	tatctatctt	tatacatata	tttaaacttt	1860
aatctacgaa	taatataatc	tatagtacta	caataatatc	agtgttttag	agaatcatat	1920
aaatgaacag	ttagacatgg	tctaaaggac	aattgagtat	tttgacaaca	ggactctaca	1980
gttttatctt	tttagtgtgc	atgtgttctc	ctttttttt	gcaaatagct	tcacctatat	2040
aatacttcat	ccattttatt	agtacatcca	tttagggttt	agggttaatg	gtttttatag	2100
actaatttt	ttagtacatc	tattttattc	tattttagcc	tctaaattaa	gaaaactaaa	2160
actctatttt	agtttttta	tttaataatt	tagatataaa	atagaataaa	ataaagtgac	2220
taaaaattaa	acaaataccc	tttaagaaat	taaaaaaact	aaggaaacat	ttttcttgtt	2280
tcgagtagat	aatgccagcc	tgttaaacgc	cgtcgacgag	tctaacggac	accaaccagc	2340

gaaccagcag cgtcgcgtcg ggccaagcga agcagacggc acggcatctc tgtcgctgcc 2400 2460 tetggacece tetegagagt teegeteeac egttggactt geteegetgt eggeateeag 2520 2580 acggcacggc agctacgggg gattcetttc ccaccgctcc ttcgctttcc cttcctcgcc 2640 cgccgtaata aatagacacc ccctccacac cctctttccc caacctcgtg ttgttcggag cgcacacaca cacaaccaga tctcccccaa atccacccgt cggcacctcc gcttcaaggt 2700 acgccgctcg tcctccccc cccccctct ctaccttctc tagatcggcg ttccggtcca 2760 tgcatggtta gggcccggta gttctacttc tgttcatgtt tgtgttagat ccgtgtttgt 2820 2880 gttagatccg tgctgctagc gttcgtacac ggatgcgacc tgtacgtcag acacgttctg 2940 attgctaact tgccagtgtt tctctttggg gaatcctggg atggctctag ccgttccgca 3000 gacgggatcg atttcatgat tttttttgtt tcgttgcata gggtttggtt tgcccttttc 3060 ctttatttca atatatgccg tgcacttgtt tgtcgggtca tcttttcatg cttttttttg 3120 tcttggttgt gatgatgtgg tctggttggg cggtcgttct agatcggagt agaattctgt ttcaaactac ctggtggatt tattaatttt ggatctgtat gtgtgtgcca tacatattca 3180 3240 tagttacgaa ttgaagatga tggatggaaa tatcgatcta ggataggtat acatgttgat 3300 gcgggtttta ctgatgcata tacagagatg ctttttgttc gcttggttgt gatgatgtgg 3360 tgtggttggg cggtcgttca ttcgttctag atcggagtag aatactgttt caaactacct 3420 ggtgtattta ttaattttgg aactgtatgt gtgtgtcata catcttcata gttacgagtt taagatggat ggaaatatcg atctaggata ggtatacatg ttgatgtggg ttttactgat 3480 3540 gcatatacat gatggcatat gcagcatcta ttcatatgct ctaaccttga gtacctatct 3600 attataataa acaagtatgt tttataatta ttttgatctt gatatacttg gatgatggca tatgcagcag ctatatgtgg atttttttag ccctgccttc atacgctatt tatttgcttg 3660 3720 gtactgtttc ttttgtcgat gctcaccctg ttgtttggtg ttacttctgc agggtacccc 3780 cggggtcgac catggccaac aagcacctga gcctctccct cttcctcgtg ctcctcggcc 3840 tetecgeete eetegeeage ggegaaattg tgeteaegea gteteeagge accetgtett 3900 tgtctccagg ggaaaaagcc accctctcct gcagggccag tcagagtgtt agtagcgcct acttagectg gtaccageag aaacetggee aggeteecag geteeteate tatggtgeat 3960 4020 ccagcagggc cactggcatc ccagacaggt tcagtggcag tgggtctggg acagacttca 4080 ctctcaccat cagcagactg gaacctgaag attttgcagt gtattactgt cagcagtatg

4140 gtaggtcacc cactttcggc ggagggacca aggtggagat caaacgaact gtggctgcac 4200 catctgtctt catcttcccg ccatctgatg agcagttgaa atctggaact gcctctgttg 4260 tgtgcctgct gaataacttc tatcccagag aggccaaagt acagtggaag gtggataacg 4320 ccctccaatc gggtaactcc caggagagtg tcacagagca ggacagcaag gacagcacct 4380 acagcctcag caacaccctg acgctgagca aagcagacta cgagaaacac aaagtctacg 4440 cctgcgaagt cacccatcag ggcctgagat cgcccgtcac aaagagcttc aacaggggag agtgttgagt taaactgagg gcactgaagt cgcttgatgt gctgaattgt ttgtgatgtt 4500 4560 ggtggcgtat tttgtttaaa taagtaagca tggctgtgat tttatcatat gatcgatctt 4620 tggggtttta tttaacacat tgtaaaatgt gtatctatta ataactcaat gtataagatg tgttcattct tcggttgcca tagatctgct tatttgacct gtgatgtttt gactccaaaa 4680 4740 accaaaatca caactcaata aactcatgga atatgtccac ctgtttcttg aagagttcat 4800 ctaccattcc agttggcatt tatcagtgtt gcagcggcgc tgtgctttgt aacataacaa 4860 ttgttacggc atatatccaa cggccggcct aggccacggt ggccagatcc actagttcta gagcggccgc ttaattaaat ttaaatgttt aaactaggcc tcctgcaggg tttaaacttg 4920 4980 ccgtggccta ttttcagaag aagttcccaa tagtagtcca aaatttttgt aacgaaggga 5040 gcataatagt tacatgcaaa ggaaaactgc cattctttag aggggatgct tgtttaagaa 5100 caaaaaatat atcactttct tttgttccaa gtcattgcgt atttttttaa aaatatttgt 5160 tccttcgtat atttcgagct tcaatcactt tatggttctt tgtattctgg ctttgctgta 5220 aatcgtagct aaccttcttc ctagcagaaa ttattaatac ttgggatatt tttttagaat 5280 caagtaaatt acatattacc accacatcga gctgctttta aattcatatt acagccatat 5340 aggettgatt cattttgcaa aatttccagg atattgacaa egttaactta ataatatett gaaatattaa agctattatg attaggggtg caaatggacc gagttggttc ggtttatatc 5400 5460 aaaatcaaac caaaccaact atatcggttt ggattggttc ggttttgccg ggttttcagc attttctggt ttttttttg ttagatgaat attattttaa tcttactttg tcaaattttt 5520 5580 5640 aatattctta taggagaatt ttcttaataa cacatgatat ttatttattt tagtcgtttg actaattttt cgttgatgta cactttcaaa gttaaccaaa tttagtaatt aagtataaaa 5700 atcaatatga tacctaaata atgatatgtt ctatttaatt ttaaattatc gaaatttcac 5760

5820 ttcaaattcg aaaaagatat ataagaattt tgatagattt tgacatatga atatggaaga 5880 acaaagagat tgacgcattt tagtaacact tgataagaaa gtgatcgtac aaccaattat 5940 ttaaagttaa taaaaatgga gcacttcata tttaacgaaa tattacatgc cagaagagtc 6000 gcaaatattt ctagatattt tttaaagaaa attctataaa aagtcttaaa ggcatatata 6060 taaaaactat atatttatat tttggtttgg ttcgaatttg ttttactcaa taccaaacta aattagacca aatataattg ggatttttaa tcgcggccca ctagtcaccg gtgtgcttgg 6120 cgtaatcatg gtcatagctg tttcctgtgt gaaattgtta tccgctcaca attccacaca 6180 acatacgage eggaageata aagtgtaaag eetggggtge etaatgagtg agetaactea 6240 cattaattgc gttgcgctca ctgcccgctt tccagtcggg aaacctgtcg tgccagctgc 6300 6360 attaatgaat cggccaacgc gcggggagag gcggtttgcg tattgggcgc tcttccgctt 6420 cctcgctcac tgactcgctg cgctcggtcg ttcggctgcg gcgagcggta tcagctcact 6480 caaaggcggt aatacggtta tccacagaat caggggataa cgcaggaaag aacatgtgag 6540 caaaaggcca gcaaaaggcc aggaaccgta aaaaggccgc gttgctggcg tttttccata 6600 ggctccgccc ccctgacgag catcacaaaa atcgacgctc aagtcagagg tggcgaaacc 6660 cgacaggact ataaagatac caggegtttc cccctggaag ctccctcgtg cgctctcctg ttccgaccct gccgcttacc ggatacctgt ccgcctttct cccttcggga agcgtggcgc 6720 6780 tttctcatag ctcacgctgt aggtatctca gttcggtgta ggtcgttcgc tccaagctgg 6840 gctgtgtgca cgaacccccc gttcagcccg accgctgcgc cttatccggt aactatcgtc 6900 ttgagtccaa cccggtaaga cacgacttat cgccactggc agcagccact ggtaacagga 6960 ttagcagagc gaggtatgta ggcggtgcta cagagttctt gaagtggtgg cctaactacg 7020 gctacactag aaggacagta tttggtatct gcgctctgct gaagccagtt accttcggaa 7080 aaagagttgg tagctcttga tccggcaaac aaaccaccgc tggtagcggt ggtttttttg 7140 tttgcaagca gcagattacg cgcagaaaaa aaggatctca agaagatcct ttgatctttt ctacggggtc tgacgctcag tggaacgaaa actcacgtta agggattttg gtcatgagat 7200 7260 tatcaaaaag gatcttcacc tagatccttt taaattaaaa atgaagtttt aaatcaatct 7320 aaagtatata tgagtaaact tggtctgaca gttaccaatg cttaatcagt gaggcaccta 7380 tctcagcgat ctgtctattt cgttcatcca tagttgcctg actccccgtc gtgtagataa 7440 ctacgatacg ggagggctta ccatctggcc ccagtgctgc aatgataccg cgagacccac gctcaccggc tccagattta tcagcaataa accagccagc cggaagggcc gagcgcagaa 7500

gtggtcctgc	aactttatcc	gcctccatcc	agtctattaa	ttgttgccgg	gaagctagag	7560
taagtagttc	gccagttaat	agtttgcgca	acgttgttgc	cattgctaca	ggcatcgtgg	7620
tgtcacgctc	gtcgtttggt	atggcttcat	tcagctccgg	ttcccaacga	tcaaggcgag	7680
ttacatgatc	ccccatgttg	tgcaaaaaag	cggttagctc	cttcggtcct	ccgatcgttg	7740
tcagaagtaa	gttggccgca	gtgttatcac	tcatggttat	ggcagcactg	cataattctc	7800
ttactgtcat	gccatccgta	agatgctttt	ctgtgactgg	tgagtactca	accaagtcat	7860
tctgagaata	gtgtatgcgg	cgaccgagtt	gctcttgccc	ggcgtcaata	cgggataata	7920
ccgcgccaca	tagcagaact	ttaaaagtgc	tcatcattgg	aaaacgttct	tcggggcgaa	7980
aactctcaag	gatcttaccg	ctgttgagat	ccagttcgat	gtaacccact	cgtgcaccca	8040
actgatcttc	agcatctttt	actttcacca	gcgtttctgg	gtgagcaaaa	acaggaaggc	8100
aaaatgccgc	aaaaaggga	ataagggcga	cacggaaatg	ttgaatactc	atactcttcc	8160
tttttcaata	ttattgaagc	atttatcagg	gttattgtct	catgagcgga	tacatatttg	8220
aatgtattta	gaaaaataaa	caaatagggg	ttccgcgcac	atttccccga	aaagtgccac	8280
ctgacgtcta	agaaaccatt	attatcatga	cattaaccta	taaaaatagg	cgtatcacga	8340
ggccctttcg	tc					8352
<210× 17						

<210> 17

<211> 12380

<212> DNA

<213> Artificial sequence

<220>

<223> pDAB637 (ubi H+L) sequence

<400> 17

60 tegegegttt eggtgatgae ggtgaaaace tetgacacat geageteeeg gagaeggtea cagcttgtct gtaagcggat gccgggagca gacaagcccg tcaggggcgcg tcagcgggtg 120 180 ttggcgggtg tcggggctgg cttaactatg cggcatcaga gcagattgta ctgagagtgc 240 accatatgcg gtgtgaaata ccgcacagat gcgtaaggag aaaataccgc atcaggcgcc 300 attegecatt caggetgege aactgttggg aagggegate ggtgegggee tettegetat 360 tacgccagct ggcgaaaggg ggatgtgctg caaggcgatt aagttgggta acgccagggt tttcccagtc acgacgttgt aaaacgacgg ccagtgaatt acaccggtgt gatcatgggc 420 cgcgattaaa aatcccaatt atatttggtc taatttagtt tggtattgag taaaacaaat 480

tcgaaccaaa ccaaaatata aatatatagt ttttatatat atgcctttaa gactttttat 540 600 agaattttct ttaaaaaata tctagaaata tttgcgactc ttctggcatg taatatttcg 660 ttaaatatqa aqtqctccat ttttattaac tttaaataat tggttgtacg atcactttct 720 tatcaagtgt tactaaaatg cgtcaatctc tttgttcttc catattcata tgtcaaaatc 780 tatcaaaatt cttatatatc tttttcgaat ttgaagtgaa atttcgataa tttaaaatta aatagaacat atcattattt aggtatcata ttgattttta tacttaatta ctaaatttgg 840 ttaactttga aagtgtacat caacgaaaaa ttagtcaaac gactaaaata aataaatatc 900 atgtgttatt aagaaaattc tcctataaga atattttaat agatcatatg tttgtaaaaa 960 aaattaattt ttactaacac atatatttac ttatcaaaaa tttgacaaag taagattaaa 1020 1080 ataatattca tctaacaaaa aaaaaaccag aaaatgctga aaacccggca aaaccgaacc 1140 aatccaaacc gatatagttg gtttggtttg attttgatat aaaccgaacc aactcggtcc 1200 atttgcaccc ctaatcataa tagctttaat atttcaagat attattaagt taacgttgtc 1260 aatatcctgg aaattttgca aaatgaatca agcctatatg gctgtaatat gaatttaaaa 1320 gcagctcgat gtggtggtaa tatgtaattt acttgattct aaaaaaatat cccaagtatt 1380 aataatttct qctaqqaaqa aggttagcta cgatttacag caaagccaga atacaaagaa 1440 ccataaagtg attgaagctc gaaatatacg aaggaacaaa tatttttaaa aaaatacgca 1500 atgacttgga acaaaagaaa gtgatatatt ttttgttctt aaacaagcat cccctctaaa 1560 gaatggcagt tttcctttgc atgtaactat tatgctccct tcgttacaaa aattttggac 1620 tactattggg aacttettet gaaaatagtg gecacegett aattaacace ggtggeeegg 1680 ccqcattccc gggaagctag gccaccgtgg cccgcctgca ggggaagctt gcatgcctgc 1740 agateceegg ggatecteta gagtegaeet geagtgeage gtgaeeeggt egtgeeeete tctagagata atgagcattg catgtctaag ttataaaaaa ttaccacata ttttttttgt 1800 cacacttgtt tgaagtgcag tttatctatc tttatacata tatttaaact ttaatctacg 1860 1920 aataatataa totatagtao tacaataata toagtgtttt agagaatoat ataaatgaac 1980 agttagacat ggtctaaagg acaattgagt attttgacaa caggactcta cagttttatc 2040 tttttagtgt gcatgtgttc tcctttttt ttgcaaatag cttcacctat ataatacttc 2100 atccatttta ttagtacatc catttagggt ttagggttaa tggtttttat agactaattt ttttagtaca tctattttat tctattttag cctctaaatt aagaaaacta aaactctatt 2160 2220 ttagtttttt tatttaataa tttagatata aaatagaata aaataaagtg actaaaaaatt

2280 aaacaaatac cctttaagaa attaaaaaaa ctaaggaaac atttttcttg tttcgagtag 2340 ataatgccag cctgttaaac gccgtcgacg agtctaacgg acaccaacca gcgaaccagc 2400 agcgtcgcgt cgggccaagc gaagcagacg gcacggcatc tctgtcgctg cctctggacc 2460 cetetegaga gtteegetee accgttggae ttgeteeget gteggeatee agaaattgeg 2520 tggcggagcg gcagacgtga gccggcacgg caggcggcct cctcctcctc tcacggcacg geagetaegg gggatteett teecaceget eettegettt eeetteeteg eeegeegtaa 2580 taaatagaca ccccctccac accctctttc cccaacctcg tgttgttcgg agcgcacaca 2640 cacacaacca gatetecece aaatecaece gteggeacet eegetteaag gtaegeeget 2700 eqtecteece ecceecect etetacette tetagategg egtteeggte eatgeatggt 2760 2820 tagggcccgg tagttctact tctgttcatg tttgtgttag atccgtgttt gtgttagatc 2880 cgtgctgcta gcgttcgtac acggatgcga cctgtacgtc agacacgttc tgattgctaa 2940 cttgccagtg tttctctttg gggaatcctg ggatggctct agccgttccg cagacgggat 3000 cgatttcatg atttttttg tttcgttgca tagggtttgg tttgcccttt tcctttattt 3060 caatatatgc cgtgcacttg tttgtcgggt catcttttca tgctttttt tgtcttggtt 3120 gtgatgatgt ggtctggttg ggcggtcgtt ctagatcgga gtagaattct gtttcaaact 3180 acctggtgga tttattaatt ttggatctgt atgtgtgtgc catacatatt catagttacg 3240 aattgaagat gatggatgga aatatcgatc taggataggt atacatgttg atgcgggttt 3300 tactgatgca tatacagaga tgctttttgt tcgcttggtt gtgatgatgt ggtgtggttg 3360 ggcggtcgtt cattcgttct agatcggagt agaatactgt ttcaaactac ctggtgtatt 3420 tattaatttt ggaactgtat gtgtgtgtca tacatcttca tagttacgag tttaagatgg 3480 atggaaatat cgatctagga taggtataca tgttgatgtg ggttttactg atgcatatac atgatggcat atgcagcatc tattcatatg ctctaacctt gagtacctat ctattataat 3540 3600 aaacaagtat gttttataat tattttgatc ttgatatact tggatgatgg catatgcagc 3660 agctatatgt ggattttttt agccctgcct tcatacgcta tttatttgct tggtactgtt tettttgteg atgeteacce tgttgtttgg tgttaettet geagggtaec eeeggggteg 3720 accatggcca acaagcacct gagcetetee etetteeteg tgeteetegg ceteteegee 3780 teeetegeea geggeeaggt teagetegtg eagteagggg etgaggtgaa gaageetggg 3840 3900 tecteggtga aggteteetg caaggettet ggaggtteet teageageta tgetateaae

3960 tgggtgcgac aggcccctgg acaagggctt gagtggatgg gagggctcat gcctatcttt 4020 gggacaacaa actacgcaca gaagttccag gacagactca cgattaccgc ggacgtatcc 4080 acgagtacag cctacatgca gctgagcggc ctgacatatg aagacacggc catgtattac 4140 tgtgcgagag ttgcctatat gttggaacct accgtcactg cagggggttt ggacgtctgg 4200 ggcaaaggga ccacggtcac cgtctcccca gcatccccga ccagccccaa ggtcttcccg ctgagectet geageaceea geeagatggg aacgtggtea tegeetgeet ggteeaggge 4260 4320 ttcttccccc aggagccact cagtgtgacc tggagcgaaa gcggacaggg cgtgaccgcc agaaacttcc cacccagcca ggatgcctcc ggggacctgt acaccacgag cagccagctg 4380 accetgeegg ceacaeagtg cetageegge aagteegtga catgeeaegt gaageaetae 4440 4500 acgaatecca gecaggatgt gaetgtgeee tgeecagtte ceteaactee acetaceeea 4560 tctccctcaa ctccacctac cccatctccc tcatgctgcc acccccgact gtcactgcac 4620 cgaccggccc tcgaggacct gctcttaggt tcagaagcga acctcacgtg cacactgacc 4680 ggcctgagag atgcctcagg tgtcaccttc acctggacgc cctcaagtgg gaagagcgct gttcaaggac cacctgagcg tgacctctgt ggctgctaca gcgtgtccag tgtcctgccg 4740 4800 ggctgtgccg agccttggaa tcatgggaag accttcactt gcactgctgc ctaccccgag 4860 tccaagaccc cgctaaccgc caccctctca aaatccggaa acacattccg gcccgaggtc 4920 cacctgctgc cgccgccgtc ggaggagctg gccctgaacg agctggtgac gctgacgtgc 4980 ctggcacgtg gcttcagccc caaggacgtg ctggttcgct ggctgcaggg gtcacaggag 5040 ctgccccgcg agaagtacct gacttgggca tcccggcagg agcccagcca gggcaccacc 5100 accttcgctg tgaccagcat actgcgcgtg gcagccgagg actggaagaa gggggacacc 5160 ttctcctgca tggtgggcca cgaggccctg ccgctggcct tcacacagaa gaccatcgac cgcttggcgg gtaaacccac ccatgtcaat gtgtctgttg tcatggcgga ggtggacggc 5220 5280 acctgctact gagttaaact gagggcactg aagtcgcttg atgtgctgaa ttgtttgtga tgttggtggc gtattttgtt taaataagta agcatggctg tgattttatc atatgatcga 5340 5400 tctttggggt tttatttaac acattgtaaa atgtgtatct attaataact caatgtataa 5460 gatgtgttca ttcttcggtt gccatagatc tgcttatttg acctgtgatg ttttgactcc 5520 aaaaaccaaa atcacaactc aataaactca tggaatatgt ccacctgttt cttgaagagt 5580 tcatctacca ttccagttgg catttatcag tgttgcagcg gcgctgtgct ttgtaacata 5640 acaattgtta cggcatatat ccaacggccg gcctaggcca cggtggccag atccactagt

5700 tctagagcgg ccgcgggcaa attcccggga agctaggcca ccgtggcccg cctgcagggg aagettgeat geetgeagat eeceggggat eetetagagt egacetgeag tgeagegtga 5760 5820 cccggtcgtg cccctctcta gagataatga gcattgcatg tctaagttat aaaaaattac 5880 cacatatttt ttttgtcaca cttgtttgaa gtgcagttta tctatcttta tacatatatt taaactttaa tctacgaata atataatcta tagtactaca ataatatcag tgttttagag 5940 aatcatataa atgaacagtt agacatggtc taaaggacaa ttgagtattt tgacaacagg 6000 actctacagt tttatctttt tagtgtgcat gtgttctcct ttttttttgc aaatagcttc 6060 6120 acctatataa tacttcatcc attttattag tacatccatt tagggtttag ggttaatggt 6180 ttttatagac taattttttt agtacatcta ttttattcta ttttagcctc taaattaaga 6240 aaactaaaac tctattttag tttttttatt taataattta gatataaaat agaataaaat 6300 aaagtgacta aaaattaaac aaataccctt taagaaatta aaaaaactaa ggaaacattt 6360 ttcttgtttc gagtagataa tgccagcctg ttaaacgccg tcgacgagtc taacggacac 6420 caaccagcga accagcagcg tcgcgtcggg ccaagcgaag cagacggcac ggcatctctg 6480 tegetgeete tggaeceete tegagagtte egeteeaceg ttggaettge teegetgteg 6540 gcatccagaa attgcgtggc ggagcggcag acgtgagccg gcacggcagg cggcctcctc 6600 ctcctctcac ggcacggcag ctacggggga ttcctttccc accgctcctt cgctttccct 6660 tectegeeeg eegtaataaa tagacaeeee eteeacaeee tettteeeea acetegtgtt 6720 gttcggagcg cacacacaca caaccagatc tcccccaaat ccacccgtcg gcacctccgc 6780 ttcaaggtac gccgctcgtc ctccccccc cccctctct accttctcta gatcggcgtt ccggtccatg catggttagg gcccggtagt tctacttctg ttcatgtttg tgttagatcc 6840 gtgtttgtgt tagatccgtg ctgctagcgt tcgtacacgg atgcgacctg tacgtcagac 6900 6960 acgttctgat tgctaacttg ccagtgtttc tctttgggga atcctgggat ggctctagcc 7020 7080 cccttttcct ttatttcaat atatgccgtg cacttgtttg tcgggtcatc ttttcatgct 7140 ttttttttgtc ttggttgtga tgatgtggtc tggttgggcg gtcgttctag atcggagtag 7200 aattotgttt caaactacct ggtggattta ttaattttgg atctgtatgt gtgtgccata 7260 catattcata gttacgaatt gaagatgatg gatggaaata tcgatctagg ataggtatac 7320 atgttgatgc gggttttact gatgcatata cagagatgct ttttgttcgc ttggttgtga

7380 tgatgtggtg tggttgggcg gtcgttcatt cgttctagat cggagtagaa tactgtttca 7440 aactacctgg tgtatttatt aattttggaa ctgtatgtgt gtgtcataca tcttcatagt 7500 tacgagttta agatggatgg aaatatcgat ctaggatagg tatacatgtt gatgtgggtt 7560 ttactgatgc atatacatga tggcatatgc agcatctatt catatgctct aaccttgagt 7620 acctatctat tataataaac aagtatgttt tataattatt ttgatcttga tatacttgga tgatggcata tgcagcagct atatgtggat ttttttagcc ctgccttcat acgctattta 7680 tttgcttggt actgtttctt ttgtcgatgc tcaccctgtt gtttggtgtt acttctgcag 7740 7800 ggtacccccg gggtcgacca tggccaacaa gcacctgagc ctctccctct tcctcgtgct cctcggcctc tccgcctccc tcgccagcgg cgaaattgtg ctcacgcagt ctccaggcac 7860 7920 cctgtctttg tctccagggg aaaaagccac cctctcctgc agggccagtc agagtgttag 7980 tagegeetae ttageetggt accageagaa acetggeeag geteecagge teeteateta 8040 tggtgcatcc agcagggcca ctggcatccc agacaggttc agtggcagtg ggtctgggac 8100 agacttcact ctcaccatca gcagactgga acctgaagat tttgcagtgt attactgtca gcagtatggt aggtcaccca ctttcggcgg agggaccaag gtggagatca aacgaactgt 8160 8220 ggctgcacca tctgtcttca tcttcccgcc atctgatgag cagttgaaat ctggaactgc 8280 ctctgttgtg tgcctgctga ataacttcta tcccagagag gccaaagtac agtggaaggt 8340 ggataacgcc ctccaatcgg gtaactccca ggagagtgtc acagagcagg acagcaagga 8400 cagcacctac agcctcagca acaccctgac gctgagcaaa gcagactacg agaaacacaa 8460 agtctacgcc tgcgaagtca cccatcaggg cctgagatcg cccgtcacaa agagcttcaa caggggagag tgttgagtta aactgagggc actgaagtcg cttgatgtgc tgaattgttt 8520 gtgatgttgg tggcgtattt tgtttaaata agtaagcatg gctgtgattt tatcatatga 8580 tcgatctttg gggttttatt taacacattg taaaatgtgt atctattaat aactcaatgt 8640 8700 ataagatgtg ttcattcttc ggttgccata gatctgctta tttgacctgt gatgttttga 8760 ctccaaaaac caaaatcaca actcaataaa ctcatggaat atgtccacct gtttcttgaa 8820 gagttcatct accattccag ttggcattta tcagtgttgc agcggcgctg tgctttgtaa 8880 cataacaatt gttacggcat atatccaacg gccggcctag gccacggtgg ccagatccac 8940 tagttctaga gcggccgctt aattaaattt aaatgtttaa actaggcctc ctgcagggtt 9000 taaacttgcc gtggcctatt ttcagaagaa gttcccaata gtagtccaaa atttttgtaa 9060 cgaagggagc ataatagtta catgcaaagg aaaactgcca ttctttagag gggatgcttg

9120 tttaaqaaca aaaaatatat cactttcttt tqttccaagt cattgcgtat tttttaaaa 9180 atatttgttc cttcgtatat ttcgagcttc aatcacttta tggttctttg tattctggct 9240 ttgctgtaaa tcgtagctaa ccttcttcct agcagaaatt attaatactt gggatatttt 9300 tttagaatca agtaaattac atattaccac cacatcgagc tgcttttaaa ttcatattac agccatatag gcttgattca ttttgcaaaa tttccaggat attgacaacg ttaacttaat 9360 aatatcttga aatattaaag ctattatgat taggggtgca aatggaccga gttggttcgg 9420 tttatatcaa aatcaaacca aaccaactat atcggtttgg attggttcgg ttttgccggg 9480 9540 ttttcagcat tttctggttt tttttttgtt agatgaatat tattttaatc ttactttgtc 9600 9660 tctattaaaa tattcttata ggagaatttt cttaataaca catgatattt atttatttta 9720 gtcgtttgac taatttttcg ttgatgtaca ctttcaaagt taaccaaatt tagtaattaa 9780 gtataaaaat caatatgata cctaaataat gatatgttct atttaatttt aaattatcga 9840 aatttcactt caaattcgaa aaagatatat aagaattttg atagattttg acatatgaat 9900 atggaagaac aaagagattg acgcatttta gtaacacttg ataagaaagt gatcgtacaa 9960 ccaattattt aaagttaata aaaatggagc acttcatatt taacgaaata ttacatgcca 10020 qaaqaqtcqc aaatatttct aqatattttt taaaqaaaat tctataaaaa gtcttaaagg 10080 10140 ccaaactaaa ttagaccaaa tataattggg atttttaatc gcggcccact agtcaccggt 10200 gtgcttggcg taatcatggt catagctgtt tcctgtgtga aattgttatc cgctcacaat 10260 tccacacaac atacgagccg gaagcataaa gtgtaaagcc tggggtgcct aatgagtgag ctaactcaca ttaattgcgt tgcgctcact gcccgctttc cagtcgggaa acctgtcgtg 10320 ccagctgcat taatgaatcg gccaacgcgc ggggagaggc ggtttgcgta ttgggcgctc 10380 10440 ttccgcttcc tcgctcactg actcgctgcg ctcggtcgtt cggctgcggc gagcggtatc 10500 ageteactea aaggeggtaa taeggttate cacagaatea ggggataaeg caggaaagaa 10560 catgtgagca aaaggccagc aaaaggccag gaaccgtaaa aaggccgcgt tgctggcgtt 10620 tttccatagg ctccgcccc ctgacgagca tcacaaaaat cgacgctcaa gtcagaggtg gcgaaacccg acaggactat aaagatacca ggcgtttccc cctggaagct ccctcgtgcg 10680 10740 ctctcctgtt ccgaccctgc cgcttaccgg atacctgtcc gcctttctcc cttcgggaag

cgtggcgctt tctcatagct cacgctgtag gtatctcagt tcggtgtagg tcgttcgctc 10860 caagetggge tgtgtgcacg aacceeecgt teageeegae egetgegeet tateeggtaa 10920 ctatcgtctt gagtccaacc cggtaagaca cgacttatcg ccactggcag cagccactgg 10980 taacaggatt agcagagcga ggtatgtagg cggtgctaca gagttcttga agtggtggcc taactacggc tacactagaa ggacagtatt tggtatctgc gctctgctga agccagttac 11040 cttcggaaaa agagttggta gctcttgatc cggcaaacaa accaccgctg gtagcggtgg 11100 tttttttgtt tgcaagcagc agattacgcg cagaaaaaaa ggatctcaag aagatccttt 11160 11220 gatcttttct acggggtctg acgctcagtg gaacgaaaac tcacgttaag ggattttggt 11280 catgagatta tcaaaaagga tcttcaccta gatcctttta aattaaaaat gaagttttaa 11340 atcaatctaa agtatatatg agtaaacttg gtctgacagt taccaatgct taatcagtga 11400 ggcacctate teagegatet gtetattteg tteatecata gttgcetgae teecegtegt 11460 gtagataact acgatacggg agggcttacc atctggcccc agtgctgcaa tgataccgcg 11520 agacccacge teaceggete cagatttate ageaataaae cagecageeg gaagggeega gcgcagaagt ggtcctgcaa ctttatccgc ctccatccag tctattaatt gttgccggga 11580 11640 agctagagta agtagttcgc cagttaatag tttgcgcaac gttgttgcca ttgctacagg 11700 categtggtg teaegetegt egtttggtat ggetteatte ageteeggtt eccaaegate 11760 aaggcgagtt acatgateee ceatgttgtg caaaaaageg gttageteet teggteetee 11820 gatcgttgtc agaagtaagt tggccgcagt gttatcactc atggttatgg cagcactgca taattctctt actgtcatgc catccgtaag atgcttttct gtgactggtg agtactcaac 11880 11940 caagtcattc tgagaatagt gtatgcggcg accgagttgc tcttgcccgg cgtcaatacg ggataatacc gcgccacata gcagaacttt aaaagtgctc atcattggaa aacgttcttc 12000 ggggcgaaaa ctctcaagga tcttaccgct gttgagatcc agttcgatgt aacccactcg 12060 12120 tgcacccaac tgatcttcag catcttttac tttcaccagc gtttctgggt gagcaaaaac 12180 aggaaggcaa aatgccgcaa aaaagggaat aagggcgaca cggaaatgtt gaatactcat 12240 actetteett ttteaatatt attgaageat ttateagggt tattgtetea tgageggata 12300 catatttgaa tgtatttaga aaaataaaca aataggggtt ccgcgcacat ttccccgaaa 12360 agtgccacct gacgtctaag aaaccattat tatcatgaca ttaacctata aaaataggcg 12380 tatcacgagg ccctttcgtc

<210> 18

<211> 16

<212> PRT

<213> Artificial sequence

<220>

<223> CDR3 region of heavy chain FabHSV 8-CDR3

<400> 18

Val Ala Tyr Met Leu Glu Pro Thr Val Thr Ala Gly Gly Leu Asp Val 1 5 10 15

<210> 19

<211> 122

<212> PRT

<213> Artificial sequence

<220>

<223> Heavy chain V region FabSHV 8

<400> 19

Leu Glu Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser Ser Val Lys
1 10 15

Val Ser Cys Lys Ala Ser Gly Gly Ser Phe Ser Ser Tyr Ala Ile Asn 20 25 30

Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met Gly Gly Leu 35 40 45

Met Pro Ile Phe Gly Thr Thr Asn Tyr Ala Gln Lys Phe Gln Asp Arg 50 55 60

Leu Thr Ile Thr Ala Asp Val Ser Thr Ser Thr Ala Tyr Met Gln Leu 70 75 80

Ser Gly Leu Thr Tyr Glu Asp Thr Ala Met Tyr Tyr Cys Ala Arg Val 85 90 95

Ala Tyr Met Leu Glu Pro Thr Val Thr Ala Gly Gly Leu Asp Val Trp
100 105 110

Gly Gln Gly Thr Thr Val Thr Val Ala Ser 115 120

<210> 20

```
<211> 18
<212> PRT
<213> Artificial sequence
<220>
<223> tryptic+ Asp-N peptide of N269
<400> 20
Asp Leu Leu Gly Ser Glu Ala Asn Leu Thr Cys Thr Leu Thr Gly
Leu Arg
<210> 21
<211> 18
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T1
<400> 21
Glu Ile Val Leu Thr Gln Ser Pro Gly Thr Leu Ser Leu Ser Pro Gly
                                   10
Glu Arg
<210> 22
<211> 6
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T2
<400> 22
Ala Thr Leu Ser Cys Arg
<210> 23
<211> 22
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T3
```

<400> 23

Ala Ser Gln Ser Val Ser Ser Ala Tyr Leu Ala Trp Tyr Gln Gln Lys 1 5 10 15

Pro Gly Gln Ala Pro Arg

<210> 24

<211> 9

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment L-T4

<400> 24

Leu Leu Ile Tyr Gly Ala Ser Ser Arg 1 5

<210> 25

<211> 7

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment L-T5

<400> 25

Ala Thr Gly Ile Pro Asp Arg

<210> 26

<211> 16

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment L-T6

<400> 26

Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Arg
1 5 10 15

<210> 27

<211> 16

<212> PRT

<213> Artificial sequence

```
<220>
<223> peptide tryptic fragment L-T7
<400> 27
Leu Glu Pro Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Gly Arg
<210> 28
<211> 9
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T8
<400> 28
Ser Pro Thr Phe Gly Gln Gly Thr Lys
<210> 29
<211> 18
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T11
<400> 29
Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln
Leu Lys
<210> 30
<211> 16
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T12
<400> 30
Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg
<210> 31
<211> 4
```

```
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T14
<400> 31
Val Gln Trp Lys
<210> 32
<211> 20
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T15
<400> 32
Val Asp Asn Ala Leu Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu
Gln Asp Ser Lys
           20
<210> 33
<211> 14
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T16
<400> 33
Asp Ser Thr Tyr Ser Leu Ser Asn Thr Leu Thr Leu Ser Lys
<210> 34
<211> 5
<212> PRT
<213> Artificial sequence
<223> peptide tryptic fragment L-T17
<400> 34
Ala Asp Tyr Glu Lys
               5
```

```
<210> 35
<211> 12
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T19
<400> 35
Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Arg
               5
<210> 36
<211> 5
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T20
<400> 36
Ser Pro Val Thr Lys
<210> 37
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T21
<400> 37
Ser Phe Asn Arg
<210> 38
<211> 23
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T5-6
<400> 38
Ala Thr Gly Ile Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp
```

Phe Thr Leu Thr Ile Ser Arg 20

<210> 39

<211> 32

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment L-T6-7

<400> 39

Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Arg
1 5 10 15

Leu Glu Pro Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Gly Arg
20 25 30

<210> 40

<211> 13

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment L-T8-9

<400> 40

Ser Pro Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys 1 5 10

<210> 41

<211> 19

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment L-T10-11

<400> 41

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu 1 5 10 15

Gln Leu Lys

<210> 42

<211> 19

<212> PRT

```
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T12-13
<400> 42
Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg
Glu Ala Lys
<210> 43
<211> 7
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T13-14
<400> 43
Glu Ala Lys Val Gln Trp Lys
<210> 44
<211> 24
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T14-15
<400> 44
Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser Gln Glu
               5
Ser Val Thr Glu Gln Asp Ser Lys
<210> 45
<211>
<212> PRT
<213> Artificial sequence
<223> peptide tryptic fragment L-T17-18
```

<400> 45

```
Ala Asp Tyr Glu Lys His Lys
               5
1
<210> 46
<211> 14
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T18-19
<400> 46
His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Arg
<210> 47
<211> 9
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T20-21
<400> 47
Ser Pro Val Thr Lys Ser Phe Asn Arg
               5
<210> 48
<211> 7
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment L-T21-22
<400> 48
Ser Phe Asn Arg Gly Glu Cys
<210> 49
<211> 12
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T1
<400> 49
Glu Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys
```

1 10 <210> 50 <211> 7 <212> PRT <213> Artificial sequence <220> <223> peptide tryptic fragment H-T2 <400> 50 Lys Pro Gly Ser Ser Val Lys 5 <210> 51 <211> 15 <212> PRT <213> Artificial sequence <220> <223> peptide tryptic fragment H-T4 <400> 51 Ala Ser Gly Gly Ser Phe Ser Ser Tyr Ala Ile Asn Trp Val Arg 15 10 <210> 52 <211> 25 <212> PRT <213> Artificial sequence <223> peptide tryptic fragment H-T5 <400> 52 Gln Ala Pro Gly Gln Gly Leu Glu Trp Met Gly Gly Leu Met Pro Ile 5 10 Phe Gly Thr Thr Asn Tyr Ala Gln Lys <210> 53 <211> 4 <212> PRT <213> Artificial sequence <220>

<223> peptide tryptic fragment H-T6

<400> 53 Phe Gln Asp Arg <210> 54 <211> 31 <212> PRT <213> Artificial sequence <220> <223> peptide tryptic fragment H-T7 <400> 54 Leu Thr Ile Thr Ala Asp Val Ser Thr Ser Thr Ala Tyr Met Gln Leu 10 Ser Gly Leu Thr Tyr Glu Asp Thr Ala Met Tyr Tyr Cys Ala Arg 20 25 <210> 55 <211> 34 <212> PRT <213> Artificial sequence <220> <223> peptide tryptic fragment H-T8 <400> 55 Val Ala Tyr Met Leu Glu Pro Thr Val Thr Ala Gly Gly Leu Asp Val Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Pro Thr Ser 25 Pro Lys <210> 56 <211> 44 <212> PRT <213> Artificial sequence <220> <223> peptide tryptic fragment H-T9 <400> 56

Val Phe Pro Leu Ser Leu Cys Ser Thr Gln Pro Asp Gly Asn Val Val

-47-

10

15

Thr Trp Ser Glu Ser Gly Gln Gly Val Thr Ala Arg 35 40

<210> 57

1

<211> 30

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment H-T10

5

<400> 57

Asn Phe Pro Pro Ser Gln Asp Ala Ser Gly Asp Leu Tyr Thr Thr Ser $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Ser Gln Leu Thr Leu Pro Ala Thr Gln Cys Leu Ala Gly Lys 20 25 30

<210> 58

<211> 7

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment H-T11

<400> 58

Ser Val Thr Cys His Val Lys 1 5

<210> 59

<211> 38

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment H-T12

<400> 59

His Tyr Thr Asn Pro Ser Gln Asp Val Thr Val Pro Cys Pro Val Pro 1 5 10 15

Ser Thr Pro Pro Thr Pro Ser Pro Ser Thr Pro Pro Thr Pro Ser Pro 20 25 30

Ser Cys Cys His Pro Arg 35

<210> 60

<211> 27

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment H-T13

<400> 60

Leu Ser Leu His Arg Pro Ala Leu Glu Asp Leu Leu Gly Ser Glu
1 5 10 15

Ala Asn Leu Thr Cys Thr Leu Thr Gly Leu Arg
20 25

<210> 61

<211> 15

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment H-T14

<400> 61

Asp Ala Ser Gly Val Thr Phe Thr Trp Thr Pro Ser Ser Gly Lys 1 5 10 15

<210> 62

<211> 9

<212> PRT

<213> Artificial sequence

<220>

<223> peptide tryptic fragment H-T15

<400> 62

Ser Ala Val Gln Gly Pro Pro Glu Arg 1 5

<210> 63

<211> 23

<212> PRT

```
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T16
<400> 63
Asp Leu Cys Gly Cys Tyr Ser Val Ser Ser Val Leu Pro Gly Cys Ala
Glu Pro Trp Asn His Gly Lys
<210> 64
<211> 12
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T17
<400> 64
Thr Phe Thr Cys Thr Ala Ala Tyr Pro Glu Ser Lys
<210> 65
<211> 9
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T18
<400> 65
Thr Pro Leu Thr Ala Thr Leu Ser Lys
               5
<210> 66
<211> 32
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T19
<400> 66
Ser Gly Asn Thr Phe Arg Pro Glu Val His Leu Leu Pro Pro Pro Ser
```

10

Glu Glu Leu Ala Leu Asn Glu Leu Val Thr Leu Thr Cys Leu Ala Arg 20 25 30

```
<210> 67
<211> 5
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T20
<400> 67
Gly Phe Ser Pro Lys
<210> 68
<211> 5
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T21
<400> 68
Asp Val Leu Val Arg
               5
<210> 69
<211> 10
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T22
<400> 69
Trp Leu Gln Gly Ser Gln Glu Leu Pro Arg
<210> 70
<211> 7
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T24
<400> 70
```

Tyr Leu Thr Trp Ala Ser Arg

5 1 <210> 71 <211> 17 <212> PRT <213> Artificial sequence <220> <223> peptide tryptic fragment H-T25 <400> 71 Gln Glu Pro Ser Gln Gly Thr Thr Thr Phe Ala Val Thr Ser Ile Leu 5 10 Arg <210> 72 <211> 7 <212> PRT <213> Artificial sequence <223> peptide tryptic fragment H-T26 <400> 72 Val Ala Ala Glu Asp Trp Lys <210> 73 <211> 20 <212> PRT <213> Artificial sequence <220> <223> peptide tryptic fragment H-T28 <400> 73 Gly Asp Thr Phe Ser Cys Met Val Gly His Glu Ala Leu Pro Leu Ala Phe Thr Gln Lys <210> 74 <211> 4 <212> PRT <213> Artificial sequence

```
<220>
<223> peptide tryptic fragment H-T29
<400> 74
Thr Ile Asp Arg
<210> 75
<211> 22
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T30
<400> 75
Leu Ala Gly Lys Pro Thr His Val Asn Val Ser Val Val Met Ala Glu
                                    10
Val Asp Gly Thr Cys Tyr
            20
<210> 76
<211> 19
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T1-2
<400> 76
Glu Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser
Ser Val Lys
<210> 77
<211> 11
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T2-3
<400> 77
Lys Pro Gly Ser Ser Val Lys Val Ser Cys Lys
```

5 10 1 <210> 78 <211> 19 <212> PRT <213> Artificial sequence <220> <223> peptide tryptic fragment H-T3-4 <400> 78 Val Ser Cys Lys Ala Ser Gly Gly Ser Phe Ser Ser Tyr Ala Ile Asn Trp Val Arg <210> 79 <211> 21 <212> PRT <213> Artificial sequence <220> <223> peptide tryptic fragment H-T17-18 <400> 79 Thr Phe Thr Cys Thr Ala Ala Tyr Pro Glu Ser Lys Thr Pro Leu Thr Ala Thr Leu Ser Lys 20 <210> 80 <211> 10 <212> PRT <213> Artificial sequence <220> <223> peptide tryptic fragment H-T20-21 <400> 80 Gly Phe Ser Pro Lys Asp Val Leu Val Arg <210> 81 <211> 15 <212> PRT <213> Artificial sequence

```
<220>
<223> peptide tryptic fragment H-T21-22
<400> 81
Asp Val Leu Val Arg Trp Leu Gln Gly Ser Gln Glu Leu Pro Arg
<210> 82
<211> 12
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T22-23
<400> 82
Trp Leu Gln Gly Ser Gln Glu Leu Pro Arg Glu Lys
<210> 83
<211> 21
<212> PRT
<213> Artificial sequence
<220>
<223> peptide tryptic fragment H-T27-28
<400> 83
Lys Gly Asp Thr Phe Ser Cys Met Val Gly His Glu Ala Leu Pro Leu
Ala Phe Thr Gln Lys
          20
<210> 84
<211>
      5118
<212> DNA
<213> Artificial sequence
<220>
<223> pDAB3014 sequence
<400> 84
                                                                60
ttagctcact cattaggcac cccaggcttt acactttatg cttccggctc gtatgttgtg
tggaattgtg agcggataac aatttcacac aggaaacagc tatgaccatg attacgccaa
                                                               180
```

240 gcttcccggg aatgcggccg ctagctagcg gccgcattcc cgggaagcta gcggccgcat 300 tcccgggaag ctagcggccg cttcccggga agcttgggct gcaggtcaat cccattgctt 360 ttgaagcage teaacattga tetetttete gaggteatte atatgettga gaagagagte gggatagtcc aaaataaaac aaaggtaaga ttacctggtc aaaagtgaaa acatcagtta 420 480 aaaggtggta taaagtaaaa tatcggtaat aaaaggtggc ccaaagtgaa atttactctt ttctactatt ataaaaattg aggatgtttt tgtcggtact ttgatacgtc atttttgtat 540 gaattggttt ttaagtttat tcgcttttgg aaatgcatat ctgtatttga gtcgggtttt 600 660 aagttcgttt gcttttgtaa atacagaggg atttgtataa gaaatatctt taaaaaaaacc catatgctaa tttgacataa tttttgagaa aaatatatat tcaggcgaat tctcacaatg 720 780 aacaataata agattaaaat agctttcccc cgttgcagcg catgggtatt ttttctagta 840 aaaataaaag ataaacttag actcaaaaca tttacaaaaa caacccctaa agttcctaaa 900 960 ccccagtcca gccaactgga caatagtctc cacaccccc cactatcacc gtgagttgtc 1020 cgcacgcacc gcacgtctcg cagccaaaaa aaaaaaaaga aagaaaaaaa agaaaaagaa 1080 aaaacagcag gtgggtccgg gtcgtggggg ccggaaacgc gaggaggatc gcgagccagc 1140 gacgaggccg gccctccctc cgcttccaaa gaaacgcccc ccatcgccac tatatacata 1200 cececete tecteceate eccecaacee taccaceace accaceacea cetecacete 1260 etececete getgeeggae gaegeetece eecteeeet eegeegeege egegeeggta 1320 accaccccgc ccctctcctc tttcttctc cgttttttt ttccgtctcg gtctcgatct 1380 ttggccttgg tagtttgggt gggcgagagg cggcttcgtg cgcgcccaga tcggtgcgcg 1440 ggaggggggg gatctcgcgg ctggggctct cgccggcgtg gatccggccc ggatctcgcg gggaatgggg ctctcggatg tagatctgcg atccgccgtt gttgggggag atgatggggg 1500 1560 gtttaaaatt tccgccatgc taaacaagat caggaagagg ggaaaagggc actatggttt atatttttat atatttctgc tgcttcgtca ggcttagatg tgctagatct ttctttcttc 1620 1680 tttttgtggg tagaatttga atccctcagc attgttcatc ggtagttttt cttttcatga 1740 tttgtgacaa atgcagcctc gtgcggagct tttttgtagg tagaccatgg cttctccgga gaggagacca gttgagatta ggccagctac agcagctgat atggccgcgg tttgtgatat 1800 1860 cgttaaccat tacattgaga cgtctacagt gaactttagg acagagccac aaacaccaca agagtggatt gatgatctag agaggttgca agatagatac ccttggttgg ttgctgaggt 1920

1980 tqaqqqtqtt qtqqctqqta ttqcttacqc tqgqccctqq aaggctagga acgcttacga 2040 ttggacagtt gagagtactg tttacgtgtc acataggcat caaaggttgg gcctaggatc 2100 cacattgtac acacatttgc ttaagtctat ggaggcgcaa ggttttaagt ctgtggttgc 2160 tgttataggc cttccaaacg atccatctgt taggttgcat gaggctttgg gatacacagc 2220 ccggggtaca ttgcgcgcag ctggatacaa gcatggtgga tggcatgatg ttggtttttg 2280 gcaaagggat tttgagttgc cagctcctcc aaggccagtt aggccagtta cccagatctg aggtaccctg agctcggtcg cagcgtgtgc gtgtccgtcg tacgttctgg ccggccgggc 2340 2400 cttgggcgcg cgatcagaag cgttgcgttg gcgtgtgtgt gcttctggtt tgctttaatt 2460 ttaccaagtt tgtttcaagg tggatcgcgt ggtcaaggcc cgtgtgcttt aaagacccac 2520 cggcactggc agtgagtgtt gctgcttgtg taggctttgg tacgtatggg ctttatttgc 2580 ttctggatgt tgtgtactac ttgggtttgt tgaattatta tgagcagttg cgtattgtaa 2640 2700 tctttaagtg ctgaattcac tggccgtcgt tttacaacgt cgtgactggg aaaaccctgg cgttacccaa cttaatcgcc ttgcagcaca tccccctttc gccagctggc gtaatagcga 2760 2820 agaggcccgc accgatcgcc cttcccaaca gttgcgcagc ctgaatggcg aatggcgcct 2880 gatgcggtat tttctcctta cgcatctgtg cggtatttca caccgcatat ggtgcactct 2940 cagtacaatc tgctctgatg ccgcatagtt aagccagccc cgacacccgc caacacccgc 3000 tgacgcgccc tgacgggctt gtctgctccc ggcatccgct tacagacaag ctgtgaccgt 3060 ctccgggagc tgcatgtgtc agaggttttc accgtcatca ccgaaacgcg cgagacgaaa 3120 gggcctcgtg atacgcctat ttttataggt taatgtcatg ataataatgg tttcttagac 3180 qtcaggtggc acttttcggg gaaatgtgcg cggaacccct atttgtttat ttttctaaat 3240 acattcaaat atgtatccgc tcatgagaca ataaccctga taaatgcttc aataatattg 3300 aaaaaggaag agtatgagta ttcaacattt ccgtgtcgcc cttattccct tttttgcggc 3360 attttgcctt cctgtttttg ctcacccaga aacgctggtg aaagtaaaag atgctgaaga 3420 tcagttgggt gcacgagtgg gttacatcga actggatctc aacagcggta agatccttga 3480 gagttttcgc cccgaagaac gttttccaat gatgagcact tttaaagttc tgctatgtgg 3540 cgcggtatta tcccgtattg acgccgggca agagcaactc ggtcgccgca tacactattc 3600 tcagaatgac ttggttgagt actcaccagt cacagaaaag catcttacgg atggcatgac

agtaagagaa	ttatgcagtg	ctgccataac	catgagtgat	aacactgcgg	ccaacttact	3660
tctgacaacg	atcggaggac	cgaaggagct	aaccgctttt	ttgcacaaca	tgggggatca	3720
tgtaactcgc	cttgatcgtt	gggaaccgga	gctgaatgaa	gccataccaa	acgacgagcg	3780
tgacaccacg	atgcctgtag	caatggcaac	aacgttgcgc	aaactattaa	ctggcgaact	3840
acttactcta	gcttcccggc	aacaattaat	agactggatg	gaggcggata	aagttgcagg	3900
accacttctg	cgctcggccc	ttccggctgg	ctggtttatt	gctgataaat	ctggagccgg	3960
tgagcgtggg	tctcgcggta	tcattgcagc	actggggcca	gatggtaagc	cctcccgtat	4020
cgtagttatc	tacacgacgg	ggagtcaggc	aactatggat	gaacgaaata	gacagatcgc	4080
tgagataggt	gcctcactga	ttaagcattg	gtaactgtca	gaccaagttt	actcatatat	4140
actttagatt	gatttaaaac	ttcattttta	atttaaaagg	atctaggtga	agatcctttt	4200
tgataatctc	atgaccaaaa	tcccttaacg	tgagttttcg	ttccactgag	cgtcagaccc	4260
cgtagaaaag	atcaaaggat	cttcttgaga	tcctttttt	ctgcgcgtaa	tctgctgctt	4320
gcaaacaaaa	aaaccaccgc	taccagcggt	ggtttgtttg	ccggatcaag	agctaccaac	4380
tctttttccg	aaggtaactg	gcttcagcag	agcgcagata	ccaaatactg	ttcttctagt	4440
gtagccgtag	ttaggccacc	acttcaagaa	ctctgtagca	ccgcctacat	acctcgctct	4500
gctaatcctg	ttaccagtgg	ctgctgccag	tggcgataag	tcgtgtctta	ccgggttgga	4560
ctcaagacga	tagttaccgg	ataaggcgca	gcggtcgggc	tgaacggggg	gttcgtgcac	4620
acagcccagc	ttggagcgaa	cgacctacac	cgaactgaga	tacctacagc	gtgagctatg	4680
agaaagcgcc	acgcttcccg	aagggagaaa	ggcggacagg	tatccggtaa	gcggcagggt	4740
cggaacagga	gagcgcacga	gggagcttcc	agggggaaac	gcctggtatc	tttatagtcc	4800
tgtcgggttt	cgccacctct	gacttgagcg	tcgatttttg	tgatgctcgt	caggggggcg	4860
gagcctatgg	aaaaacgcca	gcaacgcggc	ctttttacgg	ttcctggcct	tttgctggcc	4920
ttttgctcac	atgttctttc	ctgcgttatc	ccctgattct	gtggataacc	gtattaccgc	4980
ctttgagtga	gctgataccg	ctcgccgcag	ccgaacgacc	gagcgcagcg	agtcagtgag	5040
cgaggaagcg	tgcgcagcgg	aagagcgccc	aatacgcaaa	ccgcctctcc	ccgcgcgttg	5100
gccgattcat	taatgcag					5118

<210> 85

<211> 13680 <212> DNA <213> Artificial sequence

<220>
<223> pDAB8505 sequence

<220>
<221> misc_feature
<222> (1)..(13680)
<223> n = a or c or g or t

<400> 85 tegegegttt eggtgatgae ggtgaaaace tetgacacat geageteeeg gagaeggtea 60 cagcttgtct gtaagcggat gccgggagca gacaagcccg tcagggcgcg tcagcgggtg 120 180 ttggcgggtg tcggggctgg cttaactatg cggcatcaga gcagattgta ctgagagtgc accatatgcg gtgtgaaata ccgcacagat gcgtaaggag aaaataccgc atcaggcgcc 240 300 attequeatt caqqetqeqe aactqttggg aagggegate ggtgegggee tettegetat 360 tacgccagct ggcgaaaggg ggatgtgctg caaggcgatt aagttgggta acgccagggt tttcccagtc acgacgttgt aaaacgacgg ccagtgaatt acaccggtgt gatcatgggc 420 480 cqcqattaaa aatcccaatt atatttggtc taatttagtt tggtattgag taaaacaaat 540 . tcgaaccaaa ccaaaatata aatatatagt ttttatatat atgcctttaa gactttttat 600 agaattttct ttaaaaaata tctagaaata tttgcgactc ttctggcatg taatatttcg 660 ttaaatatga agtgctccat ttttattaac tttaaataat tggttgtacg atcactttct 720 tatcaaqtqt tactaaaatq cqtcaatctc tttgttcttc catattcata tgtcaaaatc 780 tatcaaaatt cttatatatc tttttcqaat ttgaagtgaa atttcgataa tttaaaatta 840 aatagaacat atcattattt aggtatcata ttgattttta tacttaatta ctaaatttgg 900 ttaactttga aagtgtacat caacgaaaaa ttagtcaaac gactaaaata aataaatatc atgtgttatt aagaaaattc tcctataaga atattttaat agatcatatg tttgtaaaaa 960 1020 aaattaattt ttactaacac atatatttac ttatcaaaaa tttgacaaag taagattaaa 1080 ataatattca tctaacaaaa aaaaaaccag aaaatgctga aaacccggca aaaccgaacc 1140 aatccaaacc gatatagttg gtttggtttg attttgatat aaaccgaacc aactcggtcc 1200 atttgcaccc ctaatcataa tagctttaat atttcaagat attattaagt taacgttgtc 1260 aatatcctgg aaattttgca aaatgaatca agcctatatg gctgtaatat gaatttaaaa 1320 qcagctcgat gtggtggtaa tatgtaattt acttgattct aaaaaaatat cccaagtatt 1380 aataatttct gctaggaaga aggttagcta cgatttacag caaagccaga atacaaagaa

ccataaagtg attgaagctc gaaatatacg aaggaacaaa tatttttaaa aaaatacgca 1440 1500 atgacttgga acaaaagaaa gtgatatatt ttttgttctt aaacaagcat cccctctaaa 1560 gaatggcagt tttcctttgc atgtaactat tatgctccct tcgttacaaa aattttggac 1620 tactattggg aacttettet gaaaatagtg gecacegett aattaaggeg egecatgeee 1680 ggccgcattc ccgggaagct aggccaccgt ggcccgcctg caggggaagc ttagctgaaa caaccoggco ctaaagcact atogtatoac ctatotgaaa taagtcacgg gtttcgaacg 1740 tocacttgcg togcacggaa ttgcatgttt cttgttggaa gcatattcac gcaatctcca 1800 cacataaagg tttatgtata aacttacatt tagctcagtt taattacagt cttatttgga 1860 1920 tgcatatgta tggttctcaa tccatataag ttagagtaaa aaataagttt aaattttatc ttaattcact ccaacatata tggattgagt acaatactca tgtgcatcca aacaaactac 1980 2040 ttatattgag gtgaatttgg atagaaatta aactaactta cacactaagc caatctttac 2100 tatattaaag caccagtttc aacgatcgtc ccgcgtcaat attattaaaa aactcctaca 2160 tatgtacaaa ataaggtgaa attatgtata agtgttctgg atattggttg ttggctccat 2220 2280 attcacacaa cctaatcaat agaaaacata tgttttatta aaacaaaatt tatcatatat 2340 2400 ataactagtg caacttaata catgtgtgta ttaagatgaa taagagggta tccaaataaa 2460 aaacttgttc gcttacgtct ggatcgaaag gggttggaaa cgattaaatc tcttcctagt 2520 caaaattgaa tagaaggaga tttaatctct cccaatcccc ttcgatcatc caggtgcaac 2580 cgtataagtc ctaaagtggt gaggaacacg aaacaaccat gcattggcat gtaaagctcc 2640 aagaatttgt tgtatcctta acaactcaca gaacatcaac caaaattgca cgtcaagggt attgggtaag aaacaatcaa acaaatcctc tctgtgtgca aagaaacacg gtgagtcatg 2700 2760 ccgagatcat actcatctga tatacatgct tacagctcac aagacattac aaacaactca 2820 tattgcatta caaagatcgt ttcatgaaaa ataaaatagg ccggacagga caaaaatcct 2880 tgacgtgtaa agtaaattta caacaaaaaa aaagccatat gtcaagctaa atctaattcg ttttacgtag atcaacaacc tgtagaaggc aacaaaactg agccacgcag aagtacagaa 2940 3000 tgattccaga tgaaccatcg acgtgctacg taaagagagt gacgagtcat atacatttgg 3060 caagaaacca tgaagctgcc tacagccgtc tcggtggcat agaacacaag aaattgtgtt 3120 aattaatcaa agctataaat aacgctcgca tgcctgtgca cttctccatc accaccactg

3180 ggtcttcaga ccattagctt tatctactcc agagcgcaga agaacccgat cgacaccatg 3240 ggatggagct ggatctttct cttcctcctg tcaggagctg caggtgtcca ttgccaggtt 3300 cagctcgtgc agtcaggtgc tgaggtgaag aagcctggct cctcggtgaa ggtctcctgc aaggettetg gaggtteett cageteetat getateaact gggtgaggea ageteetgga 3360 caagggettg agtggatggg agggeteatg cetatetttg ggacaacaaa etaegegeag 3420 aagttccagg acaggctcac gattaccgcg gacgtatcca cgagtacagc ctacatgcaa 3480 ctgagcggcc tgacatatga agacacggcc atgtattact gtgcgagagt tgcctacatg 3540 3600 cttgaaccta ccgtcactgc aggtggtttg gacgtctggg gccaagggac cttggtcacc 3660 gtctcctccg catccccgac cagcccgaag gtcttcccgc tgagcctctg tagcacccag 3720 ccagatggga acgtggtcat cgcctgcctg gtccagggct tcttccctca ggagccactc 3780 agtgtgacct ggagcgaaag cggacagggc gtgaccgcca ggaacttccc acccagccag 3840 gatgcctccg gagacctgta caccacgtcc agccagctga cccttccggc cacacagtgc ctagcgggca agtccgtgac atgccacgtg aagcactaca cgaatcccag ccaggatgtg 3900 3960 actgtgccct gcccagttcc ctcaactcca cctaccccat ctccctcgac tccacctacc 4020 ceateteect catgetgeca ecceaggetg teaetgeaca ggeetgeect egaggacetg 4080 ctcttaggtt cggaagcgaa cctcacgtgc acactcaccg gcctgagaga tgcgtcaggt 4140 gtcaccttca cctggacgcc ctcaagtggt aagagcgctg ttcaaggccc acctgagcgt 4200 gacctctgtg gctgctacag cgtgtccagt gtccttccgg gctgtgccga gccttggaat 4260 catgggaaga cetteacttg caetgetgee tacceegaga geaagaceee getaacegee accetetega aateeggeaa cacatteegg eeegaggtee acetgetgee geegeegteg 4320 gaggagctgg ccctgaacga gctggtgacg ctgacgtgcc tggcgcgcgg cttcagcccc 4380 aaggacgtgc tggttcgctg gctgcagggc tcacaggagc tgcctaggga gaagtacctg 4440 acttgggcat cccggcagga gcccagccaa ggcaccacca ccttcgctgt gacctcgata 4500 ctgcgcgtgg cagccgagga ctggaagaag ggtgacacct tctcctgcat ggtgggccac 4560 4620 gaggcccttc cgctggcctt cacacagaag accatcgacc gcttggcggg taaacccacc 4680 catgtcaatg tgtctgttgt catggcggag gtggacggca cctgctactg agagctcgct 4740 gagggcactg aagtcgcttg atgtgctgaa ttgtttgtga tgttggtggc gtattttgtt 4800 taaataagta agcatggctg tgattttatc atatgatcga tctttggggt tttatttaac

4860 acattgtaaa atgtgtatct attaataact caatgtataa gatgtgttca ttcttcggtt 4920 gccatagatc tgcttatttg acctgtgatg ttttgactcc aaaaaccaaa atcacaactc 4980 aataaactca tggaatatgt ccacctgttt cttgaagagt tcatctacca ttccagttgg 5040 catttatcag tgttgcagcg gcgctgtgct ttgtaacata acaattgtta cggcatatat 5100 ccaacggccg gcctagctag gccacggtgg ccagatccac tagttctaga gcggccgggc 5160 aagcggccgc attcccggga agctaggcca ccgtggcccg cctgcagggg aagcttagct gaaacaaccc ggccctaaag cactatcgta tcacctatct gaaataagtc acgggtttcg 5220 5280 aacgtccact tgcgtcgcac ggaattgcat gtttcttgtt ggaagcatat tcacgcaatc 5340 tccacacata aaggtttatg tataaactta catttagctc agtttaatta cagtcttatt 5400 tggatgcata tgtatggttc tcaatccata taagttagag taaaaaataa gtttaaattt 5460 tatcttaatt cactccaaca tatatggatt gagtacaata ctcatgtgca tccaaacaaa 5520 ctacttatat tgaggtgaat ttggatagaa attaaactaa cttacacact aagccaatct ttactatatt aaagcaccag tttcaacgat cgtcccgcgt caatattatt aaaaaactcc 5580 5640 tacatttctt tataatcaac ccgcactctt ataatctctt ctctactact ataataagag agtttatgta caaaataagg tgaaattatg tataagtgtt ctggatattg gttgttggct 5700 5760 ccatattcac acaacctaat caatagaaaa catatgtttt attaaaacaa aatttatcat 5820 5880 gcatataact agtgcaactt aatacatgtg tgtattaaga tgaataagag ggtatccaaa 5940 taaaaaactt gttcgcttac gtctggatcg aaaggggttg gaaacgatta aatctcttcc 6000 tagtcaaaat tgaatagaag gagatttaat ctctcccaat ccccttcgat catccaggtg 6060 caaccgtata agtcctaaag tggtgaggaa cacgaaacaa ccatgcattg gcatgtaaag 6120 ctccaagaat ttgttgtatc cttaacaact cacagaacat caaccaaaat tgcacgtcaa 6180 gggtattggg taagaaacaa tcaaacaaat cctctctgtg tgcaaagaaa cacggtgagt 6240 catgccgaga tcatactcat ctgatataca tgcttacagc tcacaagaca ttacaaacaa 6300 ctcatattgc attacaaaga tcgtttcatg aaaaataaaa taggccggac aggacaaaaa 6360 tccttgacgt gtaaagtaaa tttacaacaa aaaaaaagcc atatgtcaag ctaaatctaa 6420 ttcgttttac gtagatcaac aacctgtaga aggcaacaaa actgagccac gcagaagtac 6480 agaatgatto cagatgaacc atcgacgtgc tacgtaaaga gagtgacgag tcatatacat 6540 ttggcaagaa accatgaagc tgcctacagc cgtctcggtg gcatagaaca caagaaattg

6600 tgttaattaa tcaaagctat aaataacgct cgcatgcctg tgcacttctc catcaccacc 6660 actgggtctt cagaccatta gctttatcta ctccagagcg cagaagaacc cgatcgacac 6720 catgggatgg tcctggatct ttctcttcct tctgtcagga gctgcaggtg tccactgcga 6780 gategtgete aegeagtete eaggeaceet gtetttgteg ceaggggaae gtgeeaceet 6840 ctcctgccgg gccagtcagt ccgtttccag cgcgtacctt gcctggtacc agcagaagcc 6900 tggccaaget eccaggetee teatetatgg tgegteeage agggetaetg geatteeaga 6960 ccgcttctca ggcagtgggt ctgggacaga cttcacgctc accattagca ggctggaacc 7020 tgaggatttt gcagtgtact actgtcagca gtatggtcgc tcacccacgt tcggccaggg 7080 gaccaaggtg gagatcaagc gcactgtggc tgcaccgtcg gtcttcatat tcccgccatc 7140 cgatgagcag ctgaagtctg gcactgcctc tgttgtgtgc ctgctgaata acttctatcc 7200 gagagaggcg aaggtacagt ggaaggtgga taacgccctc caatcgggta actcccaaga gtccgttaca gagcaggaca gcaaggacag cacctacagc ctcagcaaca ccttgacgct 7260 7320 gagcaaagcg gactacgaga aacacaaggt ctacgcctgc gaagtcaccc atcaaggcct 7380 gcgctcgccc gtcacaaaga gcttcaaccg gggagagtgt tgagagctcg ctgagggcac 7440 tgaagtcgct tgatgtgctg aattgtttgt gatgttggtg gcgtattttg tttaaataag 7500 taagcatggc tgtgatttta tcatatgatc gatctttggg gttttattta acacattgta 7560 aaatgtgtat ctattaataa ctcaatgtat aagatgtgtt cattcttcgg ttgccataga 7620 tctgcttatt tgacctgtga tgttttgact ccaaaaacca aaatcacaac tcaataaact 7680 catggaatat gtccacctgt ttcttgaaga gttcatctac cattccagtt ggcatttatc 7740 agtgttgcag cggcgctgtg ctttgtaaca taacaattgt tacggcatat atccaacggc 7800 cggcctagct aggccacggt ggccagatcc actagttcta gagcggccgc ttaattaaat 7860 ttaaatgttt aaactaggaa atccaagctt gggctgcagg tcaatcccat tgcttttgaa 7920 gcagctcaac attgatctct ttctcgaggt cattcatatg cttgagaaga gagtcgggat 7980 agtccaaaat aaaacaaagg taagattacc tggtcaaaag tgaaaacatc agttaaaagg 8040 tggtataagt aaaatatcgg taataaaagg tggcccaaag tgaaatttac tcttttctac 8100 tattataaaa attgaggatg ttttgtcggt actttgatac gtcatttttg tatgaattgg 8160 tttttaagtt tattcgcgat tttggaaatg catatctgta tttgagtcgg gttttaagtt 8220 cgtttgcttt tgtaaataca gagggatttg tataagaaat atctttaaaa aaaccatatg

8280 ctaattttgac ataatttttg agaaaaatat atattcaggc gaattctcac aatgaacaat 8340 aataagatta aaatagcttg ccccqttgc agcgatgggt attttttcta gtaaaataaa 8400 agataaactt agactcaaaa catttacaaa aacaacccct aaagtcctaa agcccaaagt 8460 8520 agccaactgg caaatagtct ccacaccccg gcactatcac cgtgagttgt ccgcaccacc 8580 gcacgtctcg cagccaaaaa aaaaaaaaga aagaaaaaaa agaaaaagaa aaaacagcag 8640 qtqqqtccqq qtcqtqqqqq ccqqaaaaqc qaqqaqqatc qcqaqcaqcq acqaqqccqq ccctccctcc gcttccaaag aaacgccccc catcgccact atatacatac ccccccctct 8700 8760 cotoccated occeaacect accadeacea coaceaceac etectocced etegotgeeg 8820 gacgacgect ecceetece ecteegeege egeeggtaac caceeegee eteteetett 8880 tettteteeg tittittitt egieteggie tegatetitg geetiggiag titigggigg cgagagcggc ttcgtcgccc agatcggtgc gcgggagggg cgggatctcg cggctggcgt 8940 ctccgggcgt gagtcggccc ggatcctcgc ggggaatggg gctctcggat gtagatctgc 9000 9060 gatccgccgt tgttggggga gatgatgggg ggtttaaaat ttccgccatg ctaaacaaga 9120 tcaggaagag gggaaaaggg cactatggtt tatattttta tatatttctg ctgcttcgtc aggettagat gtgctagate ttetttettt ettettttg tgggtagaat ttgaateeet 9180 9240 cagcattgtt catcggtagt ttttcttttc atgatttgtg acaaatgcag cctcgtgcgg 9300 agettttttg taggtagacc atggettete eggagaggag accagttgag attaggeeag 9360 ctacagcagc tgatatggcc gcggtttgtg atatcgttaa ccattacatt gagacgtcta 9420 cagtgaactt taggacagag ccacaaacac cacaagagtg gattgatgat ctagagaggt 9480 tgcaagatag ataccettgg ttggttgctg aggttgaggg tgttgtggct ggtattgctt 9540 acgctgggcc ctggaaggct aggaacgctt acgattggac agttgagagt actgtttacg tgtcacatag gcatcaaagg ttgggcctag gatccacatt gtacacacat ttgcttaagt 9600 9660 ctatggaggc gcaaggtttt aagtctgtgg ttgctgttat aggccttcca aacgatccat 9720 ctgttaggtt gcatgaggct ttgggataca cagcccgggg tacattgcgc gcagctggat 9780 acaagcatgg tggatggcat gatgttggtt tttggcaaag ggattttgag ttgccagctc 9840 ctccaaggcc agttaggcca gttacccaga tctgaggtac caatgagctc ggtcgcagcg 9900 tgtgcgtgtc cgtcgtacgt tctggccggc cgggccttgg gcgcgcgatc agaancgttg 9960 cgttggcgtg tgtgtgcttc tggtttgctt taattttacc aagtttgttt caaggtggat

10020 cgcgtggtca aggcccgtgt gctttaaana cccaccggca ctggcagtga gtgttgctgc ttgtgtaggc tttggtacgt atgggcttta tttgcttctg gatgttgtgt actacttggg 10080 10140 tttgttgaat tattatganc agttgcgtat tgtaattcag ctgggctacc tggacattgt 10200 tatgtattaa taaatgcttt gctttcttct aaagatcttt aagtgctgaa ttcatatttc 10260 ctcctgcagg gtttaaactt gccgtggcct attttcagaa gaagttccca atagtagtcc 10320 aaaatttttg taacgaaggg agcataatag ttacatgcaa aggaaaactg ccattcttta gaggggatgc ttgtttaaga acaaaaaata tatcactttc ttttgttcca agtcattgcg 10380 10440 tattttttta aaaatatttg ttccttcgta tatttcgagc ttcaatcact ttatggttct ttgtattctg gctttgctgt aaatcgtagc taaccttctt cctagcagaa attattaata 10500 10560 cttgggatat ttttttagaa tcaagtaaat tacatattac caccacatcg agctgctttt 10620 aaattcatat tacagccata taggcttgat tcattttgca aaatttccag gatattgaca 10680 acgttaactt aataatatct tgaaatatta aagctattat gattaggggt gcaaatggac 10740 cgagttggtt cggtttatat caaaatcaaa ccaaaccaac tatatcggtt tggattggtt 10800 10860 atcttacttt gtcaaatttt tgataagtaa atatatgtgt tagtaaaaat taatttttt 10920 tacaaacata tgatctatta aaatattctt ataggagaat tttcttaata acacatgata 10980 tttatttatt ttagtcgttt gactaatttt tcgttgatgt acactttcaa agttaaccaa 11040 atttagtaat taagtataaa aatcaatatg atacctaaat aatgatatgt tctatttaat 11100 tttaaattat cgaaatttca cttcaaattc gaaaaagata tataagaatt ttgatagatt 11160 ttgacatatg aatatggaag aacaaagaga ttgacgcatt ttagtaacac ttgataagaa 11220 agtgatcgta caaccaatta tttaaagtta ataaaaatgg agcacttcat atttaacgaa 11280 atattacatg ccagaagagt cgcaaatatt tctagatatt ttttaaaagaa aattctataa aaagtcttaa aggcatatat ataaaaacta tatatttata ttttggtttg gttcgaattt 11340 11400 gttttactca ataccaaact aaattagacc aaatataatt gggattttta atcgcggccc 11460 actagtcacc ggtgtgcttg gcgtaatcat ggtcatagct gtttcctgtg tgaaattgtt 11520 atcogctcac aattocacac aacatacgag coggaagcat aaagtgtaaa gootggggtg cctaatgagt gagctaactc acattaattg cgttgcgctc actgcccgct ttccagtcgg 11580 gaaacctgtc gtgccagctg cattaatgaa tcggccaacg cgcggggaga ggcggtttgc 11640 A. 4 5

11700 gtattgggcg ctcttccgct gcgcacgctg cgcacgctgc gcacgcttcc tcgctcactg 11760 actogotgog ctoggtogtt cggctgoggc gagcggtatc agctcactca aaggcggtaa 11820 tacggttatc cacagaatca ggggataacg caggaaagaa catgtgagca aaaggccagc 11880 aaaaggccag gaaccgtaaa aaggccgcgt tgctggcgtt tttccatagg ctccgccccc 11940 ctgacgagca tcacaaaaat cgacgctcaa gtcagaggtg gcgaaacccg acaggactat 12000 aaagatacca ggcgtttccc cctggaagct ccctcgtgcg ctctcctgtt ccgaccctgc 12060 cgcttaccgg atacctgtcc gcctttctcc cttcgggaag cgtggcgctt tctcatagct cacgctgtag gtatctcagt tcggtgtagg tcgttcgctc caagctgggc tgtgtgcacg 12120 12180 aaccccccgt tcagcccgac cgctgcgcct tatccggtaa ctatcgtctt gagtccaacc 12240 cggtaagaca cgacttatcg ccactggcag cagccactgg taacaggatt agcagagcga 12300 ggtatgtagg cggtgctaca gagttcttga agtggtggcc taactacggc tacactagaa 12360 ggacagtatt tggtatctgc gctctgctga agccagttac cttcggaaaa agagttggta gctcttgatc cggcaaacaa accaccgctg gtagcggtgg tttttttgtt tgcaagcagc 12420 12480 agattacgcg cagaaaaaaa ggatctcaag aagatccttt gatcttttct acggggtctg 12540 acgctcagtg gaacgaaaac tcacgttaag ggattttggt catgagatta tcaaaaagga tcttcaccta gatcctttta aattaaaaat gaagttttaa atcaatctaa agtatatatg 12600 12660 agtaaacttg gtctgacagt taccaatgct taatcagtga ggcacctatc tcagcgatct 12720 gtctatttcg ttcatccata gttgcctgac tccccgtcgt gtagataact acgatacggg 12780 agggettace atetggeece agtgetgeaa tgatacegeg agaceeaege teaceggete 12840 cagatttatc agcaataaac cagccagccg gaagggccga gcgcagaagt ggtcctgcaa 12900 ctttatccgc ctccatccag tctattaatt gttgccggga agctagagta agtagttcgc 12960 cagttaatag tttgcgcaac gttgttgcca ttgctacagg catcgtggtg tcacgctcgt 13020 cgtttggtat ggcttcattc agctccggtt cccaacgatc aaggcgagtt acatgatccc 13080 ccatgttgtg caaaaaagcg gttagctcct tcggtcctcc gatcgttgtc agaagtaagt tggccgcagt gttatcactc atggttatgg cagcactgca taattctctt actgtcatgc 13140 13200 catccqtaaq atgcttttct gtgactggtg agtactcaac caagtcattc tgagaatagt 13260 qtatqcqqcq accqaqttqc tcttqcccqq cqtcaatacq qgataatacc qcqccacata 13320 gcagaacttt aaaagtgctc atcattggaa aacgttcttc ggggcgaaaa ctctcaagga 13380 tcttaccgct gttgagatcc agttcgatgt aacccactcg tgcacccaac tgatcttcag

catctttac tttcaccage gtttctgggt gagcaaaaac aggaaggcaa aatgccgcaa 13440
aaaagggaat aagggcgaca cggaaatgtt gaatactcat actcttcctt tttcaatatt 13500
attgaagcat ttatcagggt tattgtctca tgagcggata catatttgaa tgtatttaga 13560
aaaataaaca aataggggtt ccgcgcacat ttccccgaaa agtgccacct gacgtctaag 13620
aaaccattat tatcatgaca ttaacctata aaaataggcg tatcacgagg ccctttcgtc 13680